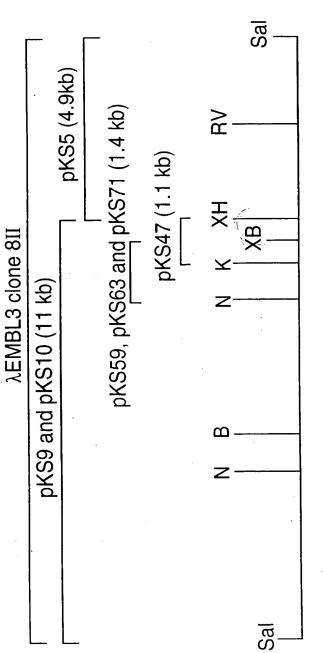


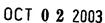
Subclones of portions of the 200 kDa protein gene from λ EMBL3 clone 8II and PCR amplification of 5' region



ATG GTG :amplified 0.7 kb

<u>FIG. 1</u>





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FIG.2A

M. catarrhalis strain 4223 AFMBL3 clone 200kDa gene

9 480 540 aaatcccaca gcacgggggg ggggtagctg tgctacaggg caagttggca 660 atcagaatgg tgatgctata tgatgatgcc tacgagttga tttgggttaa tcactctatg 300 360 420 aatactgttg ccatcattac cataatttag taacgcattt agtaacgcat ttgtaaaaat 180 cattgogccc ctttatgtgt atcatatgaa tagaatatta tgattgtatc tgattattgt 240 120 9 tgaatgacga teccaateae eagatteatt eaagtgatgt gtttgtatae geaceattta datgaatcac atctataaag tcatctttaa caaagccaca ggcacattta tggcagtggc ccctaattat ttcaatcaaa tgcctatgtc agcatgtatc atttttttaa ggtaaaccac ttttgaaact aatctattga cttaaatcac catatggtta taatttagca taatggtagg ctttttgtaa aaatcacatc gcaatattgt tctactgtta ctaccatgct catatetgta egatttgaca tgtgatatga tttaacatgt gacatgattt aacattgttt ccatggatat gggcaggtgt gctcgcctgc cgtatgatgg cgatgacacc ccatttgccc agagtacgcc atttgatata



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ggt Gly atc Ile (gtgtatgcac tctgagcttt gcccgtattg ccgcgctcgc tgtcctc gtg Val

FIG.2B

764 cat His agg Lys 캺 acc acc , daa gat a 1 Lys Lys Asp Th 15 gct tat gct caa aaa Ala Tyr Ala Gln Lys agt Ser ggc Gly acg ctc agt of Thr Leu Ser (

812 860 aag Lys 35 Sag Gln Ala gaa aat gct aac Glu Asn Ala Asn Thr 99C G1y tca Ser cgc Arg aga Arg ggt Gly 45 att Ile Pro g caa aac cag (Gln Asn Gln E gct a att Ile gcc Ala cga (Arg Glu gaa Asp. gat G1yggt Gly . Ile att gac Asp atc Ile gog Ala

50

40

908 aat gga Asn Gly gtc Val 65 Thr act agt aat aaa Ser Asn Lys' Asn Lys agt Ser 9 ggt Gly atc gcc Ala atc Ile gcc Ala Glu caa ggt G1yggc Gly

926 atc Ile Ser Ser gag Gln Caa ggt Gly agg The grades gct Ala gat Asp ggt acc (Gly Thr.) ata Ile aag Lys gat Asp Leu agt Ser ago

STENT & TRACE

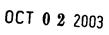
RECEIVED

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1004	1052	1100	1148	1196	1244
ggt ggt gat gta aag gct agt ggt gat gcc tcg att gcc atc Gly Gly Asp Val Lys Ala Ser Gly Asp Ala Ser Ile Ala Ile 90	gat gac tta cat ttg ctt gat cag cat ggt aat cct aaa cat 1052 Asp Asp Leu His Leu Leu Asp Gln His Gly Asn Pro Lys His 105	ggt act ctg att aac gat ctt att aac ggc cat gca gta tta 1100 Gly Thr Leu Ile Asn Asp Leu Ile Asn Gly His Ala Val Leu 120	ata cga agc tca aag gat aat gat gta aaa tat aga cgc aca Ile Arg Ser Ser Lys Asp Asn Asp Val Lys Tyr Arg Arg Thr 135	agc gga cac gcc agt act gca gtg gga gcc atg tca tat gca Ser Gly His Ala Ser Thr Ala Val Gly Ala Met Ser Tyr Ala 150	cat ttt tcc aac gcc ttt ggt aca cgg gca aca gct aaa agt His Phe Ser Asn Ala Phe Gly Thr Arg Ala Thr Ala Lys Ser 170
gcc atc Ala Ile 85	ggt agt Gly Ser 100	ccg aaa Pro Lys	aaa ga'a Lys Glu	acc gca Thr Ala	cag ggt Gln Gly 165

FIG. 2



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O S AND TO SELECT BY THE SELEC

1292	1340	1388	1436	1484	1532
gcc tat tcc ttg gca gtg ggt ctt gcc gcc aca gcc gag ggc caa tct Ala Tyr Ser Leu Ala Val Gly Leu Ala Ala Thr Ala Glu Gly Gln Ser 180 185 185	aca atc gct att ggt tct gat gca aca tct agc tcg ttg gga gcg ata Thr Ile Ala Ile Gly Ser Asp Ala Thr Ser Ser Ser Leu Gly Ala Ile 200	gcc ctt ggt gca ggt act cgt gct cag cta cag ggc agt att gcc cta Ala Leu Gly Ala Gly Thr Arg Ala Gln Leu Gln Gly Ser Ile Ala Leu 225	ggt caa ggt tct gtt gtc act cag agt gat aat aat tct aga ccg gcc Gly Gln Gly Ser Val Val Thr Gln Ser Asp Asn Asn Ser Arg Pro Ala 230	tat aca cca aat acc cag gca cta gac ccc aag ttt caa gcc acc aat Tyr Thr Pro Asn Thr Gln Ala Leu Asp Pro Lys Phe Gln Ala Thr Asn 245	aat acg aag gcg ggt cca ctt tcc att ggt agt aac tct atc aaa cgt Asn Thr Lys Ala Gly Pro Leu Ser Ile Gly Ser Asn Ser Ile Lys Arg 260 275



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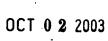
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aaa ctt gct aaa act tta aac aat ctt act gag gtg aat aca act aca Lys Leu Ala Lys Thr Leu Asn Asn Leu Thr Glu Val Asn Thr Thr 360

O SE JC130 STEEL O SER TENT 8 TRIBE

1580	1628	1676	1724	1772
aaa atc atc aat gtc ggt gca ggt gtt aat aaa acc gat gcg gtc aat	gtg gca cag cta gaa gcg gtg gtg aag tgg gct aag gag cgt aga att	act ttt cag ggt gat gat aac agt act gac gta aaa ata ggt ttg gat	aat act tta act att aaa ggt ggt gca gag acc aac gca tta acc gat	aat aat atc ggt gtg gta aaa gag gct gat aat agt ggt ctg aaa gtt
Lys Ile Ile Asn Val Gly Ala Gly Val Asn Lys Thr Asp Ala Val Asn	Val Ala Gln Leu Glu Ala Val Val Lys Trp Ala Lys Glu Arg Arg Ile	Thr Phe Gln Gly Asp Asp Asn Ser Thr Asp Val Lys Ile Gly Leu Asp	Asn Thr Leu Thr Ile Lys Gly Gly Ala Glu Thr Asn Ala Leu Thr Asp	Asn Asn Ile Gly Val Val Ilys Glu Ala Asp Asn Ser Gly Leu Ilys Val
280 280	295	310	325	340





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O STILL BY THE STREET

1868	1916	1964	2012	2060	2108
tta aat gcc aca acc aca gtt aag gta ggt agt agt agt act aca	gct gaa tta ttg agt gat agt tta acc ttt acc cag ccc aat aca ggc	agt caa agc aca agc aaa acc gtc tat ggc gtt aat ggg gtg aag ttt	act aat gca gaa aca aca gca gca atc ggc act act cgt att acc	aga gat aaa att ggc ttt gct cga gat ggt gat gtt gat gaa aaa caa	gca cca tat ttg gat aaa aaa caa ctt aaa gtg ggt agt gtt gca att
Leu Asn Ala Thr Thr Thr Val Lys Val Gly Ser Ser Ser Ser Thr Thr	Ala Glu Leu Leu Ser Asp Ser Leu Thr Phe Thr Gln Pro Asn Thr Gly	Ser Gln Ser Thr Ser Lys Thr Val Tyr Gly Val Asn Gly Val Lys Phe	Thr Asn Asn Ala Glu Thr Thr Ala Ala Ile Gly Thr Thr Arg Ile Thr	Arg Asp Lys Ile Gly Phe Ala Arg Asp Gly Asp Val Asp Glu Lys Gln	Ala Pro Tyr Leu Asp Lys Lys Gln Leu Lys Val Gly Ser Val Ala Ile
375 385	390	405	420	440	465



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2156 g atc agt aat ctt Ile Ser Asn Leu 480 ggc att gat gca ggt aat aaa aag Gly Ile Asp Ala Gly Asn Lys Lys 475 gac aat g Asp Asn (470 ata Ile acc Thr

2204			
atc gaa cag ctc aaa	Glu Glu Leu	495	
gtt acc	Val Thr	7	
gat gcg			
aac	Asn	490	
agt gct	Ser		
of ago	Gly Ser		
agg		485	
•	7		

7677	2300
re get gge ate agt gte aca eet y Ala Gly Ile Ser Val Thr Pro 510	ht ggc aat gtt acc gcc cca act r Gly Asn Val Thr Ala Pro Thr
gcc gcc aag cct act tta aac gca ggc Ala Ala Lys Pro Thr Leu Asn Ala Gly 500	act gaa ata tca gtt gat gct aag agt Thr Glu Ile Ser Val Asp Ala Lys Ser

2348		
ctt aac agt gat ggc act agt	Asn Ser Asp Gly Thr	
	Ğlu	540
aaa acc	Gly.Val Lys Thr	

530

525

520

(2396		
		Asn Ser Leu Val	
**	aag ggt	Lys Gly Ser Gly	555
	aaa ttt agt gtt	Phe Ser Val	

FIG.2G



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tat cta aat gaa gtc aat cga acg gct gac Tyr Leu Asn Glu Val Asn Arg Thr Ala Asp 570 agc Ser gcc.gaa cat ttg gca ag Ala Glu His Leu Ala St 565

254(
s ggc gca gtc a Gly Ala Val 610
gcc Ala
aat Asn
aaa Lys
aca Thr 605
acg Thr
gat Asp
aaa Liys
gct Ala
gtg Val 600
acc Thr
atc Ile
gct Ala
aac Asn

2588		
t cta acg gtt gct acc aaa	Leu Thr Val Ala	625
ggt aaa	Gly Lys Asn	620
agc atc tta aaa ctc aaa	Ile Leu Lys Leu Lys	615

2636		
ggt ctg acc	Leu Thr	
gat	r GLy	_
ğ	Ţ,	64(
a gat i	n Asp	
S Ca	k G	
tt ac	: 왕	
: ggg ctt agc caa g	NY L	535
ttt	Phe (
$\mathcal{L}_{\mathcal{L}}$		
gtt	. Val	
agc	Gly Thr	_
	_	630
a gat	s Asp	
agg	Lys	

2684		
ttg act gtt aaa gat acc	Thr Val Lys Asp	የ
ac gat ggc	G1y	んちつ
ggc	lle Gly Lys Ser	עע

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2732	2780	2828	2876	2924	2972
caa atc caa gtc ggt gct aat ggc att aaa ttt act aat gtg Gln Ile Gln Val Gly Ala Asn Gly Ile Lys Phe Thr Asn Val 675	agt aat cca ggt act ggc att gca aat acc gct cgc att acc Ser Asn Pro Gly Thr Gly Ile Ala Asn Thr Ala Arg Ile Thr 680 690	aaa att ggc ttt gct ggt tct gat ggt gca gtt gat aca aac Lys Ile Gly Phe Ala Gly Ser Asp Gly Ala Val Asp Thr Asn 695	tat ctt gat caa gac aag cta caa gtt ggc aat gtt aag att Tyr Leu Asp Gln Asp Lys Leu Gln Val Gly Asn Val Lys Ile 710	act ggc att aac gca ggt ggt aaa gcc atc aca ggg ctg tcc Thr Gly Ile Asn Ala Gly Gly Lys Ala Ile Thr Gly Leu Ser 730	ctg cct agc att gcc gat caa agt agc cgc aac ata gaa ctg Leu Pro Ser Ile Ala Asp Gln Ser Ser Arg Asn Ile Glu Leu 745
gaa 1 Glu)	: ggt 1 Gly	a gat y Asp	a cct	s aac s Asn 725	cca aca Pro Thr 740
aac Asn 660	aat Asn	aga Arg	aaa Lys	acc Thr	cca Pro 740



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ctt ggc gtc aaa acc acc aaa ctg aac aaa aca Leu Gly Val Lys Thr Thr Lys Leu Asn Lys Thr 845

gat gac aat aaa aaa c Asp Asp Asn Lys Lys L 840

3020	3068	3116	3164	3212
c gct gcc agc att aat	t aat aac aac ccc att	t gcc aat ggc aat gcc	c aaa acc agt aaa gtg	t cat cta aca ggc act
n Ala Ala Ser Ile Asn	1 Asn Asn Asn Pro Ile	e Ala Asn Gly Asn Ala	n Lys Thr Ser Lys Val	e His Leu Thr Gly Thr
770	785	800	815	0
aaa gac aaa tcc aac	ttt aac cta aaa aat	gac att gtt gac ttt	cat gat acc gct aac	gat gat aca acc att
Lys Asp Lys Ser Asn	Phe Asn Leu Lys Asn	Asp Ile Val Asp Phe	His Asp Thr Ala Asn	Asp Asp Thr Thr Ile
765	780	795	810	830
s aat aca atc caa gac	s ata tta aat aca ggc	ttt gtc tcc act tat	s acc gcc aca gta acc	
7 Asn Thr Ile Gln Asp	o Ile Leu Asn Thr Gly	Phe Val Ser Thr Tyr	r Thr Ala Thr Val Thr	
760	775	790	805	
ggc Gl <u>ý</u>	gat Asp	gac Asp	acc Thr	gta Val 820

| | | |



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agg Lys

gg Asp

ಇ೦೦

agg Lys

att Ile

ctt aat Leu Asn

ggt Gly 940

Asn

Glu

GLY

Lys

Leu

Leu 935

ggc

ggt

agg

පුදු Thr

Thr

945

3308 3356 3500 3452 Asp gac aac Asn gat Asp acc Fe Thr Asp Æ Asn Ala gat ggc gtc Val agt 930 gct Ala Gln cta F 큠 gg Ser acc tct 865 aat aat Asn Asn gac Asp Asn aat aat aac Asn Asn Asn 880 gaa aat gca Ala 895 Val gtt aat Ala Asn Thr Asn gg gcc Ala පූපු 910 aat gaa aac Asn Phe Ile atc ggc GLYtt 925 Liys gat Asp aag aac Asn gac Asp agg Lys 860 Gln Thr යින Lys 875 Thr gta Val agg acc act gaa Ala acc Thr Lys ggt Gly Ala gcc aag 890 Lys 905 Asn က္ထ His agg Val Thr aac gtg aca Asn Ile acc Thr aat gtt att gtt Val Val 920 Ile ggt G1y 855 E atc Glu ctt gaa Asn Ala aat Phe Ala Lys aag 870 Ala Asp gcc Ala 885 aac Asn gct gat acc Leu Gln ggg gaa agt Ser 900



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ggt ctt aaa gcc Gly Leu Lys Ala 960 agc Ser gtt acc ttt ggc att aac acc aca Val Thr Phe Gly Ile Asn Thr Thr 955 acg Thr 950 aat ggt Asn Gly

3644	3692
ggc aaa agc acc cta aac gac ggt ggc ttg tct att aaa aac ccc act 3ly Lys Ser Thr Leu Asn Asp Gly Gly Leu Ser Ile Lys Asn Pro Thr 965	ggt agc gaa caa atc caa gtc ggt gct gat ggc gtg aag ttt gcc aag Gly Ser Glu Gln Ile Gln Val Gly Ala Asp Gly Val Lys Phe Ala Lys 980
	-

3788		
aat ggc tca ctt gat	Asn Gly Ser Leu	1025
ygc ttt act ggg act		1020
yat gaa att	Ile	1015

gtt aat aat ggt gtt gta ggt gct ggc att gat ggc aca act cgc Val Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly Thr Thr Arg 1000 1005

3836		
agg	Lys	
ggt	Gly	
gg	GLy	
gga		.040
aac	Asn Ala	~
att	Ile	
ggc	GLy	
gac	Asp	
agg	Lys	.035
agc	Ser	
cta	: Leu Ser Lys	
Cac	His	
CC	Pro	
agg	Lys	.030
agc	Ser Lys	
agg	Lys	



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		*			*
3884	3932	3980	4028	4076	4124
aag att acc aac att caa tca ggt gag att gcc caa aac agc cat gat Lys Ile Thr Asn Ile Gln Ser Gly Glu Ile Ala Gln Asn Ser His Asp 1045 1055	gct gtg aca ggc ggc aag att tat gat tta aaa acc gaa ctt gaa aac Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn 1060 1065	aaa atc agc agt act gcc aaa aca gca caa aac tca tta cac gaa ttc Iys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu His Glu Phe 1080	tca gta gca gat gaa caa ggt aat aac ttt acg gtt agt aac cct tac Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser Asn Pro Tyr 1095	tcc agt tat gac acc tca aag acc tct gat gtc atc acc ttt gca ggt Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr Phe Ala Gly 1110	gaa aac ggc att acc acc aag gta aat aaa ggt gtg gtg cgt gtg ggc Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val Arg Val Gly 1125



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4172	4220	4268	4316	4364	4412
att gac caa acc aaa ggc tta acc acg cct aag ctg acc gtg ggt aat	aat aat ggc aaa ggc att gtc att gac agc caa aat ggt caa aat acc	atc aca gga cta agc aac act cta gct aat gtt acc aat gat aaa ggt	agc gta cgc acc aca gaa cag ggc aat ata atc aaa gac gaa gac aaa	acc cgt gcc gcc agc att gtt gat gtg cta agc gca ggc ttt aac ttg	caa ggc aat ggt gaa gcg gtt gac ttt gtc tcc act tat gac acc gtc
Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr Val Gly Asn	Asn Asn Gly Lys Gly Ile Val Ile Asp Ser Gln Asn Gly Gln Asn Thr	Ile Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn Asp Lys Gly	Ser Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp Glu Asp Lys	Thr Arg Ala Ala Ser Ile Val Asp Val Leu Ser Ala Gly Phe Asn Leu	Gln Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr Asp Thr Val
1140 1155	1160	1175	1190	1205	1220



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4460	4508	4556	4604	4652	4700
aac ttt gcc gat ggc aat gcc acc gct aag gtg acc tat gat gac	aca agc aaa acc agt aaa gtg gtc tat gat gtc aat gtg gat gat aca	ácc att gaa gtt aaa gat aaa aaa ctt ggc gta aaa acc acc aca ttg	acc agt act ggc aca ggt gct aat aaa ttt gcc cta agc aat caa gct	act ggc gat gcg ctt gtc aag gcc agt gat atc gtt gct cat cta aac	acc tta tct ggc gac atc caa act gcc aaa ggg gca agc caa gcg aac
Asn Phe Ala Asp Gly Asn Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp	Thr Ser Lys Thr Ser Lys Val Val Tyr Asp Val Asn Val Asp Asp Thr	Thr Ile Glu Val Lys Asp Lys Lys Leu Gly Val Lys Thr Thr Thr Leu	Thr Ser Thr Gly Thr Gly Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala	Thr Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala His Leu Asn	Thr Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser Gln Ala Asn
1240	1265	1270	1285	1300	1320



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aac toa goa ggo tat gtg gat got gat ggo aat aag gto ato tat gac Asn Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp 1335 1346

4796	•	•
aat gat ggc aca gtt gat	Asn Asp Gly	1360
agg	Gln Ala Lys	
acc	Ser Thr Asp Asn Lys	1350

4844		
caa gcc caa i	al Ala Gln Ala Gln Thr	1375
aaa ctg gtc	Lys Leu Val Ala	
gcc aaa gac	Ala Lys Asp Lys	1370
aaa gaa gtt	Thr Lys Glu Val	
aaa acc	Lys Thr	1365

4892	•
att aac Ile Asr	1395
tca gtc Ser Val	1390
s ggc aca ttg gct caa atg o Gly Thr Leu Ala Gln Met	1385
cca gat Pro Asp	1380

4940		
aac	Asn	
gac	Asp i	.410
ggg	$\frac{1}{2}$	
aat	Asn	
atc		
ggc	Gly Ile	
caa	Glu	405
aag	s Lys	-
agg	Liys	
aat	Asn Lys	
gcc	Ala	
gat	Asp	1400
aat	Asn	()
gta	Val	
caa	Gln 1	
gaa		

	1425	1415 1420
	r Asp Asn Lys Thr Lys	Phe Val Lys Gly Leu Glu Lys Ala
4988	gct tct gat aac aaa acc aaa	ctt gaa aaa gcc



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5036	5084	5132	5180	5228	5276
aac gcc gca gta act gtg ggt gat tta aat gcc gtt gcc caa aca ccg	ctg acc ttt gca ggg gat aca ggc aca acg gct aaa aaa ctg ggc gag	act ttg acc atc aaa ggt ggg caa aca gac acc aat aag cta acc gat	aat aac atc ggt gtg gta gca ggt act gat ggc ttc act gtc aaa ctt	gcc aaa gac cta acc aat ctt aac agc gtt aat gca ggt ggc acc aaa	att gat gac aaa ggc gtg tct ttt gta gac tca agc ggt caa gcc aaa
Asn Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala Gln Thr Pro	Leu Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys Leu Gly Glu	Thr Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys Leu Thr Asp	Asn Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr Val Lys Leu	Ala Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly Gly Thr Lys	Ile Asp Asp Lys Gly Val Ser Phe Val Asp Ser Ser Gly Gln Ala Lys
1430 1435	1445	1460 1465	1480	1495	1510

FIG.20



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gca aac acc cct gtg cta agt gcc aat ggg ctg gac ctg ggt ggc aag Ala Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys 1535 1530 1525

5372 1555 gtc atc agt aat gtg ggc aaa ggc aca aaa gat acc gac gct gcc aat Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp Ala Ala Asn 1550 1545 5420 gta caa cag tta aac gaa gta cgc aac ttg ttg ggt ctt ggt aat gct Val Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu Gly Asn Ala

1570 1565 1560 5468 Ile Lys ggt aat gat aac gct gac ggc aat cag gta aac att gcc gac atc aaa Gly Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala Asp 1575 1585 5516 Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val Ile Lys Ala 1590 tca tca tct aac cgc act gtc atc aaa gca tca ggt aaa gac cca aat

5564

ggc acg gta ctt ggc ggt aaa ggt aat aac gat acc gaa aaa ctt gcc Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala 1615 1610

FIG.2R



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5612 5756 5804 5852 5660 gcc ctg ctc gcc act tat aac gcc gca ggt cag acc aac tat ttg acc Ala Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr 1655 ttc ttc cat gtc aac gat ggc aat caa gag cct gtg gta caa ggg cgt Phe Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val Gln Gly Arg aac ggc att gac tca agt gcc tca ggc aag cac tca gtg gcg ata ggt Asn Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val Ala Ile Gly 1700 Asn Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg act ggt ggt ata caa gtg ggc gtg gat aaa gac ggc aac gct aac ggc Thr Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn Ala Asn Gly 1635 Asp Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Lys 1640 1640 aac aac ccc gca gaa gcc att gac aga ata aat gaa caa ggt atc cgc gat tta agc aat gtt tgg gtc aaa acc caa aaa gat ggc agc aaa aaa 1680 1695 1630 1645 1675 1690 1625 1640 1670 1685 1620

FIG.2S



FIG.2T

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6140

ttt ggt gtg ggc aat aac atc acc gtg acc gaa agt aac tcg Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu Ser Asn Ser 1800

gat gtc | Asp Val 1

5900 5948 5996 6092 aag cac tct ggt gcc atc ggc gac cca agc act gtt aag gct gat aac Lys His Ser Gly Ala Ile Gly Asp Pro Ser Thr Val Lys Ala Asp Asn 1765 Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr 1780 1785 acc caa gca ggc aac caa tcc atc gcc atc ggt gat aac gca caa gcc Thr Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala 1735 1746 acg ggc gat caa tcc atc gcc atc ggt aca ggc aat gtg gta gca ggt Thr Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val Val Ala Gly 1750 ggt aat aac aac cag ttt acc gat gcc act caa acc ttc cag gcc aag gca gat ggt gaa gcc gcc gtt gcc ata ggc aga caa Phe Gln Ala Lys Ala Asp Gly Glu Ala Ala Val Ala Ile Gly Arg Gln 1730 1725 1720 agt tac agt gtg



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6236 6332 6428 6428	
	cgc cgt atc Arg Arg Ile 1870 gat gcg gtc Asp Ala Val
9	acg gcg gtt fhr Ala Val
62	acc aca Thr Thr
6188	tca aac tct gcc atc agt gca ggc aca cac gca ggc Ser Asn Ser Ala Ile Ser Ala Gly Thr His Ala Gly 1820



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6476 gac cat cgt atc cac caa aac gaa aat aag gcc aat gca ggg att tca Asp His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser 1910 1910

6524 tca gcg atg gcg atg gcg tcc atg cca caa gcc tac att cct ggc aga Ser Ala Met Ala Met Ala Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg 1925

6572 tcc atg gtt acc ggg ggt att gcc acc cac aac ggt caa ggt gcg gtg 1955 Ser Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln Gly Ala Val 1950 1945 1940

6620 gca gtg gga ctg tcg aag ctg tcy yar aar yyr wur wys yw al Ala Val Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp Val Phe Lys 1970 1965 1960

8999 atc aat ggt tca gcc gat acc caa ggc cat gta ggg gcg gca gtt ggt Ile Asn Gly Ser Ala Asp Thr Gln Gly His Val Gly Ala Ala Val Gly 1975 1980 1985

6723 ggt ttt cac ttt taagccataa atcgcaagat tttacttaaa aatcaatctc Phe Ala Gly Phe His

1990



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accatagttg tataaaacag catcagcatc agtcatatta ctgatgctga tgttttttat 6783

cacttaaacc attttaccgc tcaagtgatt ctctttcacc atgaccaaat cgccattgat 6843

cataggtaaa cttattgagt aaattttatc aatgtagttg ttagatatgg ttaaaattgt 6903

gccattgacc aaaaaatgac cgatttatcc cgaaaatttc tgattatgat ccgttgacct 6963

6973

FIG.2W

gcaggtcgac





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M. catarrhalis strain 4223 genomic 200kDa gene.

FIG.3A

180 240 540 atg aat cac atc tat aaa gtc atc ttt aac aaa gcc aca ggc aca ttt 589 Met Asn His Ile Tyr Lys Val Ile Phe Asn Lys Ala Thr Gly Thr Phe 1 5 15 atcagaatgg tgatgctata tgatgatgcc tacgagttga tttgggttaa tcactctatg 300 atttgatata ttttgaaact aatctattga cttaaatcac catatggtta taatttagca 360 taatggtagg ctttttgtaa aaatcacatc gcaatattgt tctactgtta ctaccatgct 420 teceaateae cagatteatt caagtgatgt gtttgtatae geaceattta 480 ccatggatat gggcaggtgt gctcgcctgc cgtatgatgg cgatgacacc ccatttgccc 60 ccctaattat ttcaatcaaa tgcctatgtc agcatgtatc atttttttaa ggtaaaccac catatctgta cgatttgaca tgtgatatga tttaacatgt gacatgattt aacattgttt aatactgttg ccatcattac cataatttag taacgcattt agtaacgcat ttgtaaaaat cattgogccc ctttatgtgt atcatatgaa tagaatatta tgattgtatc tgattattgt tgaatgacga U



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637	989	733	781	856	877
gcc aaa tcc cac agc acg ggg ggg ggt agc 6 Ala Lys Ser His Ser Thr Gly Gly Gly Ser 25	ggc agt gta tgc act ctg agc ttt gcc cgt Gly Ser Val Cys Thr Leu Ser Phe Ala Arg 40	ctc gtg atc ggt gca acg ctc agt ggc agt Leu [Val] Ile Gly Ala Thr Leu Ser Gly Ser 55	gat acc aaa cat atc gca att ggt gaa caa Asp Thr Lys His Ile Ala Ile Gly Glu Gln 75	ggc act gcc aag gcg gac ggt gat cga gcc 8 Gly Thr Ala Lys Ala Asp Gly Asp Arg Ala 90	gct aac gca cag ggc ggt caa gcc atc gcc Ala Asn Ala Gln Gly Gly Gln Ala Ile Ala 105
tac Tyr	gtt Val	gtc Val	aaa Lys 70	tca Ser	aat Asn
gag Glu	caa Gln	gct Ala	aaa Lys	cgc Arg 85	gaa Glu
gca Ala 20	999 Gly	ctc	caa Gln	aga Arg	ggt Gly
gtg Val	aca Thr 35	gcg Ala	gct Ala	CCA	att Ile
gca Ala	gct Ala	gcc Ala 50	tat Tyr	cag Gln	gct Ala
atg Yet	tgt Cys	att Ile	gct Ala 65	aac Asn	att Ile



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925 ttg gat aag ata Leu Asp Lys Ile 125 agc agt t Ser Ser I gtc aat gga s Val Asn Gly s 120 agt aat aaa act Ser Asn Lys Thr atc ggt agt a Ile Gly Ser S

973	7
gta	
gat Asp	
ggt Gly	
ggt Gly	
atc Ile 140	
atc gcc Ile Ala	
Ser	
gag	
caa Gln 135	
ggt Gly	
acg Thr	
gct Ala	
gat Asp	
acc g Thr A 130	
ggt a Gly I	
50 0	

1021	
gac tta cat Asp Leu His 160	
gcc atc ggt agt gat Ala Ile Gly Ser Asp 155	
ggt gat gcc tcg att Gly Asp Ala Ser Ile 150	
aag gct agt (Lys Ala Ser (145	

Gly Thr Leu Ile	175
His Pro Lys	170
Leu Asp Gln His Gly Asn	165
	Asp Gln His Gly Asn Pro Lys His Pro Lys Gly Thr Leu

1117		
gaa ata cga agc	Glu Ile Arg	190
cat gca gta tta	tis Ala	185
aac gat ctt att aac ggc	Asp Leu 11e Asn	180

1165		
y gat aat gat gta aaa tat aga cgc aca acc gca agc gga cac gcc	ir Ala Ser Gly His Ala	205
cgc aca ac	Arg Thr Th	
aga	Arg	200
tat	$ ext{TY}$	
ı aaa	Lys	
gta	Val	
gat	ASp	
aat	Asn	1 Q ጊ
gat	Asp	
aag	Liys	



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O GO TO THE TRACE OF THE PATENT & TRACE OF T

1213	1261	1309	1357	1405	1453
agt act gca gtg gga gcc atg tca tat gca cag ggt cat ttt tcc aac	gcc ttt ggt aca cgg gca aca gct aaa agt gcc tat tcc ttg gca gtg	ggt ctt gcc gcc aca gcc gag ggc caa tct aca atc gct att ggt tct	gat gca aca tct agc tcg ttg gga gcg ata gcc ctt ggt gca ggt act	cgt gct cag cta cag ggc agt att gcc cta ggt caa ggt tct gtt gtc	act cag agt gat aat tct aga ccg gcc tat aca cca aat acc cag
Ser Thr Ala Val Gly Ala Met Ser Tyr Ala Gln Gly His Phe Ser Asn	Ala Phe Gly Thr Arg Ala Thr Ala Lys Ser Ala Tyr Ser Leu Ala Val	Gly Leu Ala Ala Thr Ala Glu Gly Gln Ser Thr Ile Ala Ile Gly Ser	Asp Ala Thr Ser Ser Ser Leu Gly Ala Ile Ala Leu Gly Ala Gly Thr	Arg Ala Gln Ieu Gln Gly Ser Ile Ala Ieu Gly Gln Gly Ser Val Val	Thr Gln Ser Asp Asn Asn Ser Arg Pro Ala Tyr Thr Pro Asn Thr Gln
210	225	245	260	275	290



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acc aat aat acg aag gcg ggt cca Thr Asn Asn Thr Lys Ala Gly Pro 315 ttt caa gcc a Phe Gln Ala T 310 cta gac ccc aag t Leu Asp Pro Lys P gca (Ala] 305

1549
a cgt aaa atc atc aat gtc ggt s Arg Lys Ile Ile Asn Val Gly 330
atc aaa Ile Lys
c tct n Ser
agt aac Ser Asn 325
ggt ag Gly se
att Ile
Ser
ctt Leu

gat	Asp	
gat	Asp	
ggt	GLY	
gg '	GH	365
ttt		
act	Th	
att	Ile	
aga	Arg	
	Arg	360
gag	-	
	Lys	-
gct	Ala	
tgg	Trp	1
	Lys	355
gtg	Val	
gtg	Val	

1693			
att	Thr Ile Lvs		
act	Thr		
tta	<u>[</u>	} }	
s aat act tta a	Thr		380
aat	Agn	1	
ata ggt ttg gat	Agn	3 71	
ttg	֓֞֟֟֝֟֟֝֟֟֝֟֝֟֝֟	3	
ggi	1	775	
ata	-	דד	375
aaa	<u> </u>	ν Σ	
gta	17.7	Val Lys	
gac gta	((ASD	
act act dac	E	127	
agt	, ה	S P	370
ששט	}	Asn S	

1741
c ggt gtg gta e Gly Val Val 400
c ggt e Gly
aat aat atc Asn Asn Ile 395
aat Asn 395
gat Asp
a tta acc gat a Leu Thr Asp
tta Leu
gca
aac Asn 390
acc a
gag Glu
gca Ala
ggt Gly
ggt Gly 385



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1789 E Thr act Lys agg aaa ctt gct Ala Fen Lys gtt Val agg Lys ctg Gly Leu ggt Ser agt Asn aat 405 Asp gat Glu gag agg

FIG.3F

1837 캺 큠 acc Thr 430 Ala ည္ထင aat Leu Asn tta Thr gg Thr act 425 Thr aca Asn aat gtg Val Glu gag act Thr Ferr ctt Asn aat Asn aac

1885 gat Asp Seg agt F ttg Lea 445 Glu ggg gct Ala පූපූ Thr act Thr Ser agt 440 Ser agt agt Ser agt Ser ggt Gly gta Val Lys aag Val

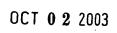
1933 agg Lys Set agc Thr Ser agc Gln Caa 460 Ser agt ggc G1Ygg Thr aat Asn Pro 455 CCC Gln gg ಇ೦೦ ttt Phe acc Thr Leu tta 450 agt

1981 aca Thr 480 Glu ggg gga Ala aat Asn Asn aat Thr act 475 Phe tt aag Lys gtg Val 999 G1y Asn aat gtt Val G1yggc $ext{TYT}$ tat Val 465

ttt Phe ggc GLYПе Lys agg Asp gat Arg aga acc Thr 490 Ile att Arg gt act-act Thr Thr Thr ggc Gly 485 atc Ile

2029

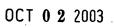
495





2077	2125	2173	2221	2269	2317
gct cga gat ggt gat gtt gat gaa aaa caa gca cca tat ttg gat aaa	aaa caa ctt aaa gtg ggt agt gtt gca att acc ata gac aat ggc att	gat gca ggt aat aaa aag atc agt aat ctt gcc aaa ggt agc agt gct	aac gat gcg gtt acc atc gaa cag ctc aaa gcc gcc aag cct act tta	aac gca ggc gct ggc atc agt gtc aca cct act gaa ata tca gtt gat	gct aag agt ggc aat gtt acc gcc cca act tac aac att ggc gtg aaa
Ala Arg Asp Gly Asp Val Asp Glu Lys Gln Ala Pro Tyr Leu Asp Lys	Lys Gln Leu Lys Val Gly Ser Val Ala Ile Thr Ile Asp Asn Gly Ile	Asp Ala Gly Asn Lys Lys Ile Ser Asn Leu Ala Lys Gly Ser Ser Ala	Asn Asp Ala Val Thr Ile Glu Gln Leu Lys Ala Ala Lys Pro Thr Leu	Asn Ala Gly Ala Gly Ile Ser Val Thr Pro Thr Glu Ile Ser Val Asp	Ala Lys Ser Gly Asn Val Thr Ala Pro Thr Tyr Asn Ile Gly Val Lys
500 500	515	530	545	575	580





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2605

ggt aaa aac ggt cta acg gtt gct acc aaa aaa gat ggt acg gtt acc Gly Lys Asn Gly Leu Thr Val Ala Thr Lys Lys Asp Gly Thr Val Thr 675

O E JC; STRATE STRATES

2365	2413	2461	2509	2557
gag ctt aac agt gat ggc act agt gat aaa ttt agt gtt aag Glu Leu Asn Ser Asp Gly Thr Ser Asp Lys Phe Ser Val Lys 595	ggt acg aac aat agc tta gtt acc gcc gaa cat ttg gca agc Gly Thr Asn Asn Ser Leu Val Thr Ala Glu His Leu Ala Ser 615	aat gaa gtc aat cga acg gct gac agt gct cta caa agc ttt Asn Glu Val Asn Arg Thr Ala Asp Ser Ala Leu Gln Ser Phe 630	aaa gaa gac gat gac gcc aac gct atc acc gtg gct Lys Glu Glu Asp Asp Asp Ala Asn Ala Ile Thr Val Ala 655	gat acg aca aaa aat gcc ggc gca gtc agc atc tta aaa ctc aaa Asp Thr Thr Lys Asn Ala Gly Ala Val Ser Ile Leu Lys Leu Lys 660
acc acc Thr Thr	ggt agt Gly Ser 610	tat cta Tyr Leu 625	acc gtt Thr Val	aaa ge Liys As



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2653	2701	2749	2797	2845	2893
ttt ggg ctt agc caa gat agc ggt ctg acc att ggc aaa agc acc cta	aac aac gat ggc ttg act gtt aaa gat acc aac gaa caa atc caa gtc	ggt gct aat ggc att aaa ttt act aat gtg aat ggt agt aat cca ggt	act ggc att gca aat acc gct cgc att acc aga gat aaa att ggc ttt	gct ggt tct gat ggt gca gtt gat aca aac aaa cct tat ctt gat caa	gac aag cta caa gtt ggc aat gtt aag att acc aac act ggc att aac
Phe Gly Leu Ser Gln Asp Ser Gly Leu Thr Ile Gly Lys Ser Thr Leu	Asn Asn Asp Gly Leu Thr Val Lys Asp Thr Asn Glu Gln Ile Gln Val	Gly Ala Asn Gly Ile Lys Phe Thr Asn Val Asn Gly Ser Asn Pro Gly	Thr Gly Ile Ala Asn Thr Ala Arg Ile Thr Arg Asp Lys Ile Gly Phe	Ala Gly Ser Asp Gly Ala Val Asp Thr Asn Lys Pro Tyr Leu Asp Gln	Asp Lys Leu Gln Val Gly Asn Val Lys Ile Thr Asn Thr Gly Ile Asn
690	705	735	740	755	770



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2947	
agc att Ser Ile 800	
aca ctg cct Thr Leu Pro	
cca Pro 795	
ctg tcc Leu Ser	
aca ggg Ihr Gly	
gcc atc a Ala Ile 7	
ggt aaa Gly Lys	
gca ggt Ala Gly 785	

atc caa gac Ile Gln Asp 815

agc cgc aac ata gaa ctg ggc aat aca Ser Arg Asn Ile Glu Leu Gly Asn Thr 805

caa agt a

gcc gat (Ala Asp (

3037	3085
gac aaa tcc aac gct gcc agc att aat gat ata tta aat aca ggc Asp Lys Ser Asn Ala Ala Ser Ile Asn Asp Ile Leu Asn Thr Gly 820	aac cta aaa aat aat aac aac ccc att gac ttt gtc tcc act tat
aaa Liys	ttt

3085			3133
ccc att gac ttt gtc tcc act tat	Lys Asn Asn Asn Pro Ile Asp Phe Val Ser	835 840	gae att oft gae ttt gee aat gge aat gee aee gee aea gta aee 3
+	 		O

31.		
aat gcc acc acc gcc aca gta acc Asn Ala Thr Thr Ala Thr Val Thr	860	
ttt gcc aat ggc Phe Ala Asn Gly	855	
gac att gtt gac Asp Ile Val Asp	850	

l Asn Val	088
sp Va	
Iyr A	
Val	875
Maj	
S A	
Thr	
S	870
Asn	,
Ala	
Thr	
Asp	4
His	865
	Asp





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TECH CENTER 1600/2900

OF JC139 JOHN STREET

3229	3277	3325	3373	3421	3469
gat gat aca acc att cat cta aca ggc act gat gac aat aaa aaa ctt	ggc gtc aaa acc acc aaa ctg aac aaa aca agt gct aat ggt aat aca	gca act aac ttt aat gtt aac tct agt gat gaa gat gcc ctt gtt aac	gcc aaa gac atc gcc gaa aat cta aac acc cta gcc aag gaa att cac	acc acc aaa ggc aca gca gac acc gcc cta caa acc ttt acc gtt aaa	aag gta gat gaa aat aat aat gct gat gac gcc aac gcc atc acc gtg
Asp Asp Thr Thr Ile His Leu Thr Gly Thr Asp Asp Asn Lys Lys Leu	Gly Val Lys Thr Thr Lys Leu Asn Lys Thr Ser Ala Asn Gly Asn Thr	Ala Thr Asn Phe Asn Val Asn Ser Ser Asp Glu Asp Ala Leu Val Asn	Ala Lys Asp Ile Ala Glu Asn Leu Asn Thr Leu Ala Lys Glu Ile His	Thr Thr Lys Gly Thr Ala Asp Thr Ala Leu Gln Thr Phe Thr Val Lys	Lys Val Asp Glu Asn Asn Asn Ala Asp Asp Ala Asn Ala Ile Thr Val
895	900	920	930	945	975



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TECH CENTER 1600/2900

ggt caa aag aac gca aat aat caa gtc aac acc cta aca ctc aaa ggt Gly Gln Lys Asn Ala Asn Asn Gln Val Asn Thr Leu Thr Leu Lys Gly 980 980

3565	
t ggt acg gtt acc ttt n Glv Thr Val Thr Phe	305
c gac aaa aat ggt e	3
ett aaa acc	116 175
gaa aac ggt ctt aat	ASII GIY LEU ASI 995

3613	
gcc ggc aaa agc acc cta aac	61y 1ys ser 111r 1.eu 1020
ctt aaa	red Lys
ca agc ggt	j (
sac acc aca	
ggc att	GIY 11e 1010

3661	
caa atc Gln Ile	1040
agc gaa Ser Glu	1035
ttg tct att aaa aac Leu Ser Ile Lys Asn	1030
gac ggt ggc ttg Asp Gly Gly Leu	1025

3709		
gtt aat aat agt gtt	sn Asn	1055
ttt gcc aag	Phe Ala Lys	
ggt gct gat ggc gtg aag	Val Gly Ala Asp Gly Val Lys	1045

3757	
acc aga gat gaa att Thr Arg Asp Glu Ile	1070
ggc aca act cgc att Gly Thr Thr Arg Ile	1065
gta ggt gct ggc att gat Val Gly Ala Gly Ile Asp	1060



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TECH CENTER 1600/290

ggc ttt act ggg act aat ggc tca ctt gat aaa agc aaa ccc cac cta Gly Phe Thr Gly Thr Asn Gly Ser Leu Asp Lys Ser Lys Pro His Leu 1075 1085

3853	
aaa aag att acc aac att caa Lys Lys Ile Thr Asn Ile Gln	1100
att aac Ile Asn	1095
agc aaa gac ggc Ser Lys Asp Gly	1090

3901	,
ggc ggc aag Gly Gly Lys 1120	
aca gr Thr G	
gtg Val	
gct Ala 1115	
gat Asp	
cat His	
agc Ser	
caa aac Gln Asn L110	
caa Gln 1110	
gcc	
att I Ile	
gag Glu	
ca ggt er Gly 105	,
tca g Ser (1105	
•	

3949	3997
att tat gat tta aaa acc gaa ctt gaa aac aaa atc agc agt act gcc Ile Tyr Asp Leu Lys Thr Glu Leu Glu Asn Lys Ile Ser Ser Thr Ala 1125	. see and the other than the the tea other day call

3997		
gaa	sp Glu Gln	20
l gta gca g	Val Ala Asp	1150
ttc tca	Ŕ	
tta cac gaa	Leu His Glu Phe S	1145
aac tca tt	Gln Asp Ser Le	
gca caa		1140
aaa aca	Lys Thr	ı

4045		
cct tac tcc agt tat gac acc tca	yr Asp	1165
gac		1160
£ £	Asn Asn Phe	1155



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ada	att 4333	. gaa 4285 . Glu	aac 4237 Asn	att 4189 Ile	ada ggc 4141 Lys Gly 1200	Thr
aat aaa ggt Asn Lys Gly acg cct aag Thr Pro Lys 1205 gac agc caa Asp Ser Gln 1220 gct aat gtt Ala Asn Val 1235	cag ggc aat ata atc aaa gac gaa gac aaa acc cgt gcc gcc agc	cta gct aat gtt acc aat gat aaa ggt agc gta cgc Leu Ala Asn Val Thr Asn Asp Lys Gly Ser Val Arg 1235	gac agc caa aat ggt caa aat acc atc aca gga cta agc Asp Ser Gln Asn Gly Gln Asn Thr Ile Thr Gly Leu Ser 1220	acc acg cct aag ctg acc gtg ggt aat aat aat Thr Thr Pro Lys Leu Thr Val Gly Asn Asn 1205	aat aaa ggt gtg gtg cgt gtg ggc att gac caa acc Asn Lys Gly Val Val Arg Val Gly Ile Asp Gln Thr 1190	Lys Thr Ser Asp Val Ile Thr Phe Ala Gly Glu Asn Gly Ile Thr 1170 1175



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TECH CENTER 1600/2900

4381	4429	4477	4525
gtt gat gtg cta agc gca ggc ttt aac ttg caa ggc aat ggt gaa gcg	gtt gac ttt gtc tcc act tat gac acc gtc aac ttt gcc gat ggc aat	gcc acc acc gct aag gtg acc tat gat gac aca agc aaa acc agt aaa	gtg gtc tat gat gtc aat gtg gat gat aca acc att gaa gtt aaa gat
Val Asp Val Leu Ser Ala Gly Phe Asn Leu Gln Gly Asn Gly Glu Ala	Val Asp Phe Val Ser Thr Tyr Asp Thr Val Asn Phe Ala Asp Gly Asn	Ala Thr Thr Ala Lys Val Thr Tyr Asp Asp Thr Ser Lys	Val Val Tyr Asp Val Asn Val Asp Asp Thr Thr Ile Glu Val Lys Asp
1265 1280	1295	1300	1315

gct aat aaa ttt gcc cta agc aat caa gct act ggc gat gcg ctt gtc Ala Asn Lys Phe Ala Leu Ser Asn Gln Ala Thr Gly Asp Ala Leu Val 1345 1350 1350 aaa aaa ctt ggc gta aaa acc acc aca ttg acc agt act ggc aca ggt Lys Lys Leu Gly Val Lys Thr Thr Thr Leu Thr Ser Thr Gly Thr Gly 1330

C 51



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TECH CENTER 1600/2900

ggc gac atc Gly Asp Ile 1375 acc tta tct g Thr Leu Ser G aag gcc agt gat atc gtt gct cat cta aac l Lys Ala Ser Asp Ile Val Ala His Leu Asn ' 1365

4717
tca gca ggc tat gtg Ser Ala Gly Tyr Val 1390
caa gcg aac aac Gln Ala Asn Asn 1385
ggg gca agc Gly Ala Ser
caa act gcc aaa Gln Thr Ala Lys 1380

4765		
gac agt acc gat aac aag tac	Asp Ser Thr Asp Asn	1405
tc atc tat	al Ile Tyr	1395 1400
gct	Asp Ala Asp Gl	

4813		
aaa acc aaa gaa gtt gcc	s Thr Lys Glu Val Ala	1420
gat ggc aca gtt gat	Asp Gly Thr Val Asp	1415
ğ	Gln Ala Liys Asr	410

4861	
aca ttg gct Thr Leu Ala	1440
ggc ac Gly Ti	ı
gat g	
Sta Pro	1435
acc Th	
caa Gln	
gcc Ala	
Cap Cap	
y gtc gcc	1430
y gtc	7
ctg	
aaa ctg	Z V
gac	Σ. Σί
aaa T-T	1425

490		
a gta aat gat gcc	n Val Asn Asp Ala 1455)))
tg aat gtc aaa tca gtc att aac aaa gaa caa gta aat gat gcc	Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu Gln	OCET 6771
caa a	Gin M	



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TECH CENTER 1600/2900

aat aaa aag caa ggc atc aat gaa gac aac gcc ttt gtt aaa gga ctt Asn Lys Lys Gln Gly Ile Asn Glu Asp Asn Ala Phe Val Lys Gly Leu 1460 1460

2002		
ğ	Ala Val Thr	1485
aac aaa acc aaa	Asn Lys Thr Lys	1480
gaa aaa gcc gct tct gat	Glu Lys Ala Ala Ser Asp	. 1475

5053		
ttt gca	Pro Leu Thr Phe Ala Gly Asp	1500
aat gcc gtt gcc caa aca	Val Ala Gln Th	1495
tt	Gly Asp Leu Asn Ala	. 1490

5101
atc aaa ggt Ile Lys Gly 1520
r ggc gag act ttg acc atc a cly Glu Thr Leu Thr Ile 1 1515
gct daa aaa ctg Ala Lys Lys Leu 1510
aca ggc aca acg Thr Gly Thr Thr 1505

5149		
s atc ggt gtg gta	ile Gly Val Val	1535
lat aac	sn Asn	
cgat a	r Asp A	1530
cta ad	Leu Th	
c aag	n Liys	
gac acc aat	Thr Asp Thr Asn Lys Leu Thr Asp Asn Asn Ile	1525
Gaa	Gly Gln Thr	
8	ਹੋ ਹੋ	

Ald Lys Asp Leu IIIL Asii 1550
rne inr val Lys Leu f 1545
Ala Gly Thr Asp Gly Pr 1540



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ctt aac agc gtt aat gca ggt ggc acc aaa att gat gac aaa ggc gtg Leu Asn Ser Val Asn Ala Gly Gly Thr Lys Ile Asp Asp Lys Gly Val 1565 1560

5293 Ser Phe Val Asp Ser Ser Gly Gln Ala Lys Ala Asn Thr Pro Val Leu 1570 tca agc ggt caa gcc aaa gca aac acc cct gtg cta tct ttt gta gac

5341 agt gcc aat ggg ctg gac ctg ggt ggc aag gtc atc agt aat gtg ggc Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val Ile Ser Asn Val Gly Ser Ala Asn Gly Leu Asp Leu Gly Gly Lys Val 1595 1590 5389 Gly Thr Lys Asp Thr Asp Ala Ala Asn Val Gln Gln Leu Asn Glu aca aaa gat acc gac gct gcc aat gta caa cag tta aac gaa 1615 1610 1605 ggc agg Liys

5437 gta cgc aac ttg ttg ggt ctt ggt aat gct ggt aat gat aac gct gac Val Arg Asn Leu Leu Gly Ieu Gly Asn Ala Gly Asn Asp Asn Ala Asp 1625 1620 5485 ggc aat cag gta aac att gcc gac atc aaa aaa gac cca aat tca ggt Gly Asn Gln Val Asn Ile Ala Asp Ile Lys Lys Asp Pro Asn Ser Gly 1640

5245



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5533	5581	5629	5677	5725	5773
tca tca tct aac cgc act gtc atc aaa gca ggc acg gta ctt ggc ggt Ser Ser Ser Asn Arg Thr Val Ile Lys Ala Gly Thr Val Leu Gly Gly 1650	aaa ggt aat aac gat acc gaa aaa ctt gcc act ggt ggt ata caa gtg Lys Gly Asn Asn Asp Thr Glu Lys Leu Ala Thr Gly Gly Ile Gln Val 1665 1675	ggc gtg gat aaa gac ggc aac gct aac ggc gat tta agc aat gtt tgg Gly Val Asp Lys Asp Gly Asn Ala Asn Gly Asp Leu Ser Asn Val Trp 1695	gtc aaa acc caa aaa gat ggc agc aaa aaa gcc ctg ctc gcc act tat Val Lys Thr Gln Lys Asp Gly Ser Lys Lys Ala Leu Leu Ala Thr Tyr 1700	aac gcc gca ggt cag acc aac tat ttg acc aac ccc gca gaa gcc Asn Ala Ala Gly Gln Thr Asn Tyr Leu Thr Asn Asn Pro Ala Glu Ala 1715	att gac aga ata aat gaa caa ggt atc cgc ttc ttc cat gtc aac gat Ile Asp Arg Ile Asn Glu Gln Gly Ile Arg Phe Phe His Val Asn Asp



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gcg ata ggt ttc cag gcc aag gca gat 5869 Ala Ile Gly Phe Gln Ala Lys Ala Asp 1770 1775	cc caa gca ggc aac caa 5917	cg ggc gat caa tcc atc 5965	ag cac tct ggt gcc atc 6013	igt tac agt gtg ggt aat 6061
	fir Gln Ala Gly Asn Gln	hr Gly Asp Gln Ser Ile	ys His Ser Gly Ala Ile	Jer Tyr Ser Val Gly Asn
	1790	1805	1820	135
one ace att ace ata ada caa ace	Val Ala Ile Gly Arg Gln Thr 1785	tcc atc gcc atc ggt gat aac gca caa gcc acg ggc g Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala Thr Gly A 1795	gcc atc ggt aca ggc aat gtg gta gca ggt aag cac tct ggt gcc atc Ala Ile Gly Thr Gly Asn Val Val Ala Gly Lys His Ser Gly Ala Ile 1810	ggc gac cca agc act gtt aag gct gat aac agt tac a Gly Asp Pro Ser Thr Val Lys Ala Asp Asn Ser Tyr 3 1825



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6109 aac aac cag ttt acc gat gcc act caa acc gat gtc ttt ggt gtg ggc Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr Asp Val Phe Gly Val Gly 1855

6157		
gcc tta ggt tca	Ala Leu	1870
gaa agt aac tog gtt	Ser Asn	1865
aat aac atc acc gtg acc	Asn Asn Ile Thr Val Thr	1860

6205		
ggc aca caa gcc aaa aaa tct	Thr Gln Ala Lys	1885
ggc aca cac gca g	Thr His Ala	1880
tct gcc atc agt gca	Ser	1875

6253		
ggt gca acc ggt acg	Gly Ala Thr Gly	1900
acc aca gas	Thr Thr Ala	
g	Gly Thr Thr	1895
ggc	Asp Gly Thr Ala Gly T	 1

6301		
gtg ggt gcc	Val Gly Ala	1920
gtt ggt gcg gtc tcc	Val Gly Ala Val Ser	1915
gct.gga caa acg gcg	Gln Thr Ala	1910
ggc ttt	31y Phe	•

634		
a ggt gag gtc agt	ਲ	1935
caa aat gtg gca gca	Gln Asn Val Ala Al	1930
tca ggt gct gaa cgc cgt atc	Ile	1075



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6445 6397 6493 6283 aac gaa aat aag gcc aat gca ggg att tca tca gcg atg gcg atg gcg Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser Ser Ala Met Ala Met Ala 1970 tcc atg cca caa gcc tac att cct ggc aga tcc atg gtt acc ggg ggt Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg Ser Met Val Thr Gly Gly 2000 att gcc acc cac aac ggt caa ggt gcg gtg gca gtg gga ctg tcg aag Ile Ala Thr His Asn Gly Gln Gly Ala Val Ala Val Gly Leu Ser Lys gcc acc agc acc gat gcg gtc aat ggt agc cag ttg tac aaa gcc acc Ala Thr Ser Thr Asp Ala Val Asn Gly Ser Gln Leu Tyr Lys Ala Thr 1940 1950 ago att goc aac goa acc aat gag ott gac oat ogt atc oac caa Gln Ser Ile Ala Asn Ala Thr Asn Glu Leu Asp His Arg Ile His Gln 1965 1995 1960 1990 1955 1985

6637 tog gat aat ggt caa tgg gta ttt aaa atc aat ggt tca gcc gat Ser Asp Asn Gly Gln Trp Val Phe Lys Ile Asn Gly Ser Ala Asp 2030 2025

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gcg gca gtt ggt gca ggt ttt cac ttt Ala Ala Val Gly Ala Gly Phe His Phe 2040 acc caa ggc cat gta ggg Thr Gln Gly His Val Gly 2035

FIG.3W

6682

6972 catcagcatc agtcatatta ctgatgctga tgttttttat cacttaaacc attttaccgc 6802 tcaagtgatt ctctttcacc atgaccaaat cgccattgat cataggtaaa cttattgagt 6862 agattttatc aatgtagttg ttagatatgg ttagaattgt gccattgacc aaaaaatgac 6922 taagocataa atogcaagat tttacttaaa aatcaatctc accatagttg tataaaacag 6742 cgatttatcc cgaaaatttc tgattatgat ccgttgacct gcaggtcgac



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M. catarrhalis strain Q8 200kDa gene

48	96	144	192	240	788
ATG aat cac atc tat aaa gtc atc ttt aac aaa gcc aca ggc aca ttt	atg gcc gtg gcg gaa tat gcc aaa tcc cac agt ac <u>g ggg ggg gg</u> t agc	tgt gct aca ggg caa gtt ggc agt gta cgc act cta agc ttt gcc cgt	att gcc gcg ctc gct gtc ctc gtg atc ggt gcg acg ctc aat ggc agt	gct tat gct caa caa att act acc aag atc gaa att ggt caa aca aac	aag ata aac aac acg ctg aaa ggc gat gcc cta gcg aca ggt gaa gca
Met Asn His Ile Tyr Lys Val Ile Phe Asn Lys Ala Thr Gly Thr Phe	Met Ala Val Ala Glu Tyr Ala Lys Ser His Ser Thr Gly Gly Gly Ser	Cys Ala Thr Gly Gln Val Gly Ser Val Arg Thr Leu Ser Phe Ala Arg	Ile Ala Ala Leu Ala Val Leu Val Ile Gly Ala Thr Leu Asn Gly Ser	Ala Tyr Ala Gln Gln Ile Thr Thr Lys Ile Glu Ile Gly Gln Thr Asn	Lys Ile Asn Asn Thr Leu Lys Gly Asp Ala Leu Ala Thr Gly Glu Ala
1 5 15	20 30	35	50 55	65 75 80	85



OCT 0 2 2003

336	384	432	480	528	576
tcc att gct ttt ggt agt ctt tct aag gca caa ggc tct caa gct att Ser Ile Ala Phe Gly Ser Leu Ser Lys Ala Gln Gly Ser Gln Ala Ile 100	gct atc ggt agt gtc aaa cca gat cct aat aat ggt agt aat ggt aat Ala Ile Gly Ser Val Lys Pro Asp Pro Asn Asn Gly Ser Asn Gly Asn 115	gta ggt tcc cac gcc aaa ggt aac gag tcc atc gcc atc ggt ggt gat Val Gly Ser His Ala Lys Gly Asn Glu Ser Ile Ala Ile Gly Gly Asp 130	gta ttg gct gag ggt gat gcc tcg att gcc atc ggt agt gat gac tta Val Leu Ala Glu Gly Asp Ala Ser Ile Ala Ile Gly Ser Asp Asp Leu 145	tat itg cct aag aat ctt gat ctg aag aat gaa itt cac aaa ctt att Tyr Leu Pro Lys Asn Leu Asp Leu Lys Asn Glu Phe His Lys Leu Ile 165	cat ggc cat gaa ata tta aaa aaa ata caa acc tca acc gat ggt aaa His Gly His Glu Ile Leu Lys Lys Ile Gln Thr Ser Thr Asp Gly Lys 180





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624	672.	720	768	816	864
atc aaa tat cga cgc aca aga gca caa ggg cac gcc agt act gca gtg	gga gcc atg tca tat gca cag ggt cat ttt tcc aac gcc ttt ggt aca	tac gca aca gct gaa gct gcc tat tcc ttg gca gta ggt ctt gcc gcc	caa gcc aca aaa caa tct tca atc gct gtt ggt tcc aat gca aaa gct	aac gcg ttt gca gcg aca gcc att ggt gga aat act gta gtt aat ttg	ggt cga ggc gtt gcc cta'ggt ttt ggt tct cag atc ctt gat agg gat
Ile Lys Tyr Arg Arg Thr Arg Ala Gln Gly His Ala Ser Thr Ala Val	Gly Ala Met Ser Tyr Ala Gln Gly His Phe Ser Asn Ala Phe Gly Thr	Tyr Ala Thr Ala Glu Ala Ala Tyr Ser Leu Ala Val Gly Leu Ala Ala	Gln Ala Thr Lys Gln Ser Ser Ile Ala Val Gly Ser Asn Ala Lys Ala	Asn Ala Phe Ala Ala Thr Ala Ile Gly Gly Asn Thr Val Val Asn Leu	Gly Arg Gly Val Ala Leu Gly Phe Gly Ser Gln Ile Leu Asp Arg Asp
200	210	225	255	260	275



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TECH CENTER 1600/2900

F acg Thr Lys agg 99t G1y 300 cta Leu පු Pro Val TYTtat gcc Ala 295 Ser agt gcc Ala ASp aat Asn 290 Asn aat

960 Ser 320 Phe Ile agg gat Asp acg Thr tct Ser 315 atc gat Asp ggt Gly Gln gg cgc Arg acc Thr 310 gcc Ala Lys Glu Cag gac Asp 305

1056 atc Ile gg gca Ala atc 11e 335 aat gtg Asn Val Lys Arg g agg Arg gtc Val gog gat agt Ser 330 acc Thr 345 Ser agc gat Asp aat Asn ogg Arg Asn aat Asn aat tct Ser ggt Gly aat Asn 325 Ser agt Asn ggt Gly aat gtc Val aat

350 geg Ala 340

Asn

Gln

ggt Gly Lys aag Phe tt Thr 365 Ile att agg Lys Arg aat cgt Asn gct Ala 360 ctg Leu gaa Glu gag Glu Leu 355 Lys Leu

Asn aat ggc GLYFer ttg 99t G1y 380 aga Arg gaa Glu gta Val Se sign aat Asn 375 Ser agc aat Asn Asn aat ggt Gly 370 gat Asp

912



OCT 0 2 2003

1200	1248	1296	1344	1392	1440
tta act att aaa ggt gat gca cag acc aac gca tta acc gaa gct aac Leu Thr Ile Lys Gly Asp Ala Gln Thr Asn Ala Leu Thr Glu Ala Asn 385 385	atc ggt gtg gta aca gat ggc aat ggt ctg aaa gtt aaa ctt gct aaa Ile Gly Val Val Thr Asp Gly Asn Gly Leu Lys Val Lys Leu Ala Lys 410	gag ctg act gga ttg acc agt gtc tcc gct acc aac aaa atc acc gtt Glu Leu Thr Gly Leu Thr Ser Val Ser Ala Thr Asn Lys Ile Thr Val 425	agt aat acc aac aac aac gcc gag cta caa agc ggt ggt ttg acc Ser Asn Thr Asn Asn Asn Ala Glu Leu Gln Ser Gly Gly Leu Thr 440	ttt agc cca ata aca ggt aca aaa aca gat aaa acc gtc tac agc att Phe Ser Pro Ile Thr Gly Thr Lys Thr Asp Lys Thr Val Tyr Ser Ile 450	gat gga ttg aag ttt act aat gat agt aat agt ata gca act aaa ggt Asp Gly Leu Lys Phe Thr Asn Asp Ser Asn Ser Ile Ala Thr Lys Gly
→ ⊢ ()	.0	0. 0			



OCT 0 2 2003

TECH CENTER 1600/2900

1728 1536 1632 1680 1584 gat Asp Arg 560 gat Asp Thr cgt Val gct gtt Val 575 aat Lys Ala Asn 495 agg acc Thr act Thr Gly Phe gga Asn act Thr Leu 510 aac Lys ggc Gly Asp Lys aat Asn 525 agg gat ggt Gly aag att Ile 540 aat Asn Glu gtc Val gct Ala gaa gtt Val Glyaca Thr 555 act Thr Phe act tt aac Asn Thr ggc gca Ala gac Asp Asn ggt Gly 570 ggt Gly 490 Leu aat ttg Ala Leu 505 tg g Ser Ala att Ile agc Ser gct gct ttt Phe GLYacc Thr Lys TYYggt agg tat ggt G1y 520 ggt Gly aat Asn Lys Pro gtc Val 535 aag cct agt Ser Glu Gaa gga Ala 550 att Lys Lys Asn agg agg aac agg Leu gtt Val Lys 565 acc Thr Śer agc 485 gag Glu Gln aat Asn Glu 500 acc Caa Asn gaa Glu gat Asp cgt Ser 515 Asn Asn 530 act Ihr gtt aat acc Thr aac ggt Gly



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TECH CENTER 1600/2900

2016

ggc

aat

agt

aac

ct

aag Lys

agg

Se ago

att Ile

aac

tat

act

සු

Thr

Asn 660

gtg Val

act Illr

acc Thr 665

Lys

GLy

Asn

Asn

Ser 670

1776 1824 1968 1872 1920 gtt Val acc Thr Ile 640 acc gg Glu att Thr IIeSet 3d ogt Arg agg att ಇ೦೦ Lys Thr 655 atc Lys Ile atc Ile Ile 9gt G1y 590 aag aat att Asn acc Cac Thr ggc GLYgtg Val His 605 aat Asn ggc GLYagg Lys gtt Val 620 gat Asp ctt F ggt Gly gog Ala 99c Gly 635 agt Ser Arg Asp gct Ala gat Ser 650 Sga gga Ala agt Glu 585 Asn TPK Asn Asp aat acc aac gat gaa Lys Ile 600 aat Asn tta Fe gtt agg att Val gat Asp GLYLen Thr cta ggt gca Ala 615 act Leu Asp Ile Ser ata cct Pro 630 gat ttg agt GlyTYTLys tat gat Asp ggt aag Gly 645 999 Pro 580 Thr aat gcc Ala CCa aga Asn aac Asn gca Ala gac aat acc Thr act Thr Asn 595 Gh Ile ctt ියුතු att Ser agg gg 610 ctc gga G1yLeu 625 aat Asn



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TECH CENTER 1600/2900

2208 2256 2064 2112 2160 2304 aat Thr acc ggt Gly Asn gat Asp 720 acc 캺 Asn Gly aac Asn aat Asn Val gg Lys 735 acg FF aag Asp Asn GLY Ile cat gat Asn Lys agc Ser agg ggc agt 750 Asn Asn His gg Gln aat Gln aat aac Gaa 685 765 gct Ala Leu 700 Asp Val Val gac Thr gtt gtc aca Ile Glyaat Lys 715 ggt att Asn TYTAsp tat agg gat Phe aaa Lys 730 Asn ggc GLyagt Seg gac Asp ttt aac Glu Ala G1ygag Phe gtt gg agc Ser ttt ggt Val 745 Leu Gly acc Thr Pro agt Ser CCB gtg Val gg ttg 760 680 Asp 695 cta gtt Val Phe Leu Thr agg Lys tt gat acc Thr Lys Ala ctc Fen ಇದ್ದ agg Lys agg gct atc 710 Ser agg Lys Asn gcc Ala ggt Gly aat aat gcc Ala agt Asn gac Asn Leu aca Thr Thr Asp aac acc 740 Ala 755 gct aac Asn acc Thr gcc ggt G1y 675 Val Asn Arg tta Feu Thr ∂ac යුරික acg Ser Ser sign 3gc 9 aat aat Ser Asn acc TPr agc gaa Glu 705



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TECH CENTER 1600/2900

aca aat ggt aat cga Thr Asn Gly Asn Arg 780 agc gat : Ser Asp ' ggt Gly gtg Val acc Thr 775 acg cct aag ctg Thr Pro Lys Leu acc Thr 770 ctc

Leu Val Ile Glu Gln 785 atc att aaa gga ttg Ile Ile Lys Gly Leu 805	Val Pro Ser Ala 790 tcc cca aca ctg Ser Pro Thr Leu	A Asp Gly Asn Ser Thr Lys Asn 795 gct agc att gcc agt cca agt 1 Pro Ser Ile Ala Ser Pro Ser 810	2448
---	--	---	------

Ser

s gaa gaa aaa gac aaa t e Glu Glu Lys Asp Lys S 830

ata gca ctg ggc aat aca atc Ile Ala Leu Gly Asn Thr Ile 820

ogc aac a

ggc Gly

2544	2592
aac cta aaa	act gtt gac
Asn Leu Lys	Thr Val Asp
aat gca ggc ttt	tcc act tat gac
Asn Ala Gly Phe	Ser Thr Tyr Asp
845	860
att gat gat gtg cta	gac aaa gac ttt gtc
Ile Asp Asp Val Leu	Asp Lys Asp Phe Val
840	855
aac gct gcc agc	aat aat ggc aaa g
Asn Ala Ala Ser	Asn Asn Gly Lys i
835	850



OCT 0 2 2003

2640	2688	2736	2784	2832	2880
ttt atc gat ggc aat gcc acc acc gcc aca gta act tat gat gaa gcc	aat caa acc agt aaa gtg gcg tat gat gtg aat gtg gat gag aaa acc	att gaa ctg aca ggc gat aat ggc aag aaa caa ctt ggc gtc aaa acc	atc aaa ctg acc gaa aca agt act aat ggt aat gca act aca ttt agt	acc gac gat gac cat gcc ctt gtt aaa gcc agt gat atc gcc ggc aat	cta aac acc cta gcc gag gaa att cac acc acc aaa ggc aca gca aac
Phe Ile Asp Gly Asn Ala Thr Thr Ala Thr Val Thr Tyr Asp Glu Ala	Asn Gln Thr Ser Lys Val Ala Tyr Asp Val Asn Val Asp Glu Lys Thr	Ile Glu Leu Thr Gly Asp Asn Gly Lys Lys Gln Leu Gly Val Lys Thr	Ile Lys Leu Thr Glu Thr Ser Thr Asn Gly Asn Ala Thr Thr Phe Ser	Thr Asp Asp Asp His Ala Leu Val Lys Ala Ser Asp Ile Ala Gly Asn	Leu Asn Thr Leu Ala Glu Glu Ile His Thr Thr Lys Gly Thr Ala Asn
865	895	900	925	930	945



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2928	2976	3024	3072	3120	3168
cta caa acc ttt acc gtt aaa aag gta gat gaa aat gat aag 2 Leu Gln Thr Phe Thr Val Lys Lys Val Asp Glu Asn Asp Lys 975	gac acc aac gcc atc acc gtg ggt aaa gat ggc aca agt ggt 2 Asp Thr Asn Ala Ile Thr Val Gly Lys Asp Gly Thr Ser Gly 980	aac acc tta aaa ctc aaa ggt aaa aac ggt ctt gat att aaa 3 Asn Thr Leu Lys Leu Lys Gly Lys Asn Gly Leu Asp Ile Lys 995	aaa gat ggt acg gtt acc ttt ggc att aac acc caa agc ggt 3 Lys Asp Gly Thr Val Thr Phe Gly Ile Asn Thr Gln Ser Gly 1015	gcc ggc gac agc act cta aac aac aat ggc ttg tct att 3 Ala Gly Asp Ser Thr Thr Leu Asn Asn Asn Gly Leu Ser Ile 1030	yct agt aac gaa caa atc caa gtc ggt gct gat ggc gtg Na Ser Asn Glu Gln Ile Gln Val Gly Ala Asp Gly Val
gcc Ala	gat Asp	gtc Val	acc gác á Thr Asp 1 1010	399 175	aaa aac acc g Lys Asn Thr 1
acc Thr	gct	aaa Lys	acc Th	ctt d Leu 1 1025	aaa Lys



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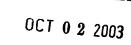
3216 ttt gcc atg gtt aat agt ggt gtt gta ggt gct ggc att gat ggc Phe Ala Met Val Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly 1060 1065 aag Lys

3264	3312
aca act cgc att acc aga gat gaa att ggc ttt act ggg act aat ggc Thr Thr Arg Ile Thr Arg Asp Glu Ile Gly Phe Thr Gly Thr Asn Gly 1075	tca ctt gat aaa agc aaa ccc cac cta agc aaa gac ggc att aac gca Ser Leu Asp Lys Ser Lys Pro His Leu Ser Lys Asp Gly Ile Asn Ala 1090

3360

ggt ggt aaa aag att acc aac att caa tca ggt gag att gcc aaa aac Gly Gly Lys Lys Ile Thr Asn Ile Gln Ser Gly Glu Ile Ala Lys Asn 1105 1110

3408	3456
agc cat gat gct gtg aca ggc ggc aag att tat gat tta aaa acc gaa Ser His Asp Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu 1125	ctt gaa aat aaa atc agc agt act gcc aaa aca gca caa aac tca tta Leu Glu Asn Lys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu 1140
(0 0)	J



TECH CENTER 1600/2900

SEP 2 6 THEN BY TRADE

3504	3552	3600	3648	3696	3744
cac gaa ttc tca gta gca gat gaa caa ggt aat aac ttt acg gtt agt	aac oot tac too agt tat gac acc toa aag acc tot gat gto atc acc	ttt gca ggt gaa aac ggc att acc acc aag gta aat aaa ggt gtg gtg	cgt gtg ggc att gac caa acc aaa ggc tta acc acg cct aag ctg acc	gtg ggt aat aat ggc aaa ggc att gtc att aac agc caa aat ggt	caa aat acc atc aca gga cta agc aac act cta gct aat gtt acc aat
His Glu Phe Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser	Asn Pro Tyr Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr	Phe Ala Gly Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val	Arg Val Gly Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr	Val Gly Asn Asn Asn Gly Lys Gly Ile Val Ile Asn Ser Gln Asn Gly	Gln Asn Thr Ile Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn
1155	1170	1185 1190	1205	1220	1235



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gat aaa ggt agc gta cgc acc aca gaa cag ggc aat ata atc aaa gac Asp Lys Gly Ser Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp 1250

384(
agc gca ggc Ser Ala Gly	1280
gtg cta a Val Leu S	
gtt gat Val Asp	1275
att Ile	
ic ago a Ser	
cgt gcc gcc Arg Ala Ala	1270
aaa acc Lys Thr	
gaa gac a Glu Asp Li	1265

ggc Gly	뀱뜠		gcc ae Ala As	ttt gcc Phe Ala	ttt gcc Phe Ala	1290 1295
aat acc acc Asn Thr Thr	ggc aat acc Gly Asn Thr	aat ggc aat acc Asn Gly Asn Thr	aat ggc aat acc Asn Gly Asn Thr	ttt gcc aat ggc aat acc Phe Ala Asn Gly Asn Thr	ttt gcc aat ggc aat acc Phe Ala Asn Gly Asn Thr	
; aat acc , Asn Thr	ggc aat Gly Asn	aat ggc aat Asn Gly Asn	aat ggc aat Asn Gly Asn	ttt gcc aat ggc aat Phe Ala Asn Gly Asn	ttt gcc aat ggc aat Phe Ala Asn Gly Asn	
: aat : Asn	ggc aat Gly Asr	aat Asn	aat Asn	ttt gcc aat Phe Ala Asn	ttt gcc aat Phe Ala Asn	
	. ggc	aat Asn	aat Asn	ttt gcc aat Phe Ala Asn	ttt gcc aat Phe Ala Asn	
ttt gcc Phe Ala	ttt Phe	gtc aac ttt Val Asn Phe	gtc aac Val Asn	gtc Val		
gcc Ala	gtc aac ttt Val Asn Phe	gtc aac Val Asn				

3984		
a gtg gtc tat gat gtc aat gtg	겂	1325
aaa acc agt aa	Lys Thr Ser Ly	
agc	Tyr Asp Asp Thr Ser	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \

4032		
aaa ctt ggc	Lys Lys Leu Gly Val Lys Thr	1340
att gaa gtt aaa gat s	ile Glu Val Lys Asp I	1335
gat gat aca acc	Asp Thr Thr	1330



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4080		
cta	Ala Leu Ser	1360
aaa ttt	Lys Phe	
gct aat	Ala Asn	1355
agg	Gly Thr Gly	
agt	Thr Ser Thr	1350
aca ttg		45
ac	딉	134

aat caa gct act ggc gat gcg ctt gtc aag gcc agt gat atc gtt gct Asn Gln Ala Thr Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Val Ala 1375

cat cta aac acc tta tct ggc gac atc caa act gcc aaa ggg gca agc His Leu Asn Thr Leu Ser Gly Asp Ile Gln Thr Ala Lys Gly Ala Ser 1380 1390

caa gog aac aac toa goa ggo tat gtg gat got gat ggo aat aag gto Gln Ala Asn Asn Ser Ala Gly Tyr Val Asp Ala Asp Gly Asn Iys Val 1395

atc tat gac agt acc gat aac aag tac tat caa gcc aaa aat gat ggc Ile Tyr Asp Ser Thr Asp Asn Lys Tyr Tyr Gln Ala Lys Asn Asp Gly 1410

aca gtt gat aaa acc aaa gaa gtt gcc aaa gac aaa ctg gtc gcc caa Thr Val Asp Lys Thr Lys Glu Val Ala Lys Asp Lys Leu Val Ala Gln

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TECH CENTER 1600/2900

gcc caa acc cca gat ggc aca ttg gct caa atg aat gtc aaa tca gtc Ala Gln Thr Pro Asp Gly Thr Leu Ala Gln Met Asn Val Iys Ser Val 1445

4416		
aag caa ggc atc aat	Lys Gln Gly Ile Asn	1470
gat gcc aat aas	n Asp Ala Asn Lys	1465
att aac aaa gaa caa gta aat	Ile Asn Lys Glu Gln Val Asn	1460

gaa gac aac gcc ttt gtt aaa gga ctt gaa aaa gcc gct tct gat aac Glu Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala Ser Asp Asn 1475 1486	4464		
gac aac gcc ttt gtt aaa gga ctt gaa aaa gcc gct tct Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala Ser 1475 1485	aac	Asn	
gac aac gcc ttt gtt aaa gga ctt gaa aaa gcc gct tct Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala Ser 1475 1485	gat	Asp	
gac aac gcc ttt gtt aaa gga ctt gaa aaa gcc gct Asp Asn Ala Phe Val Lys Gly Leu Glu Lys Ala Ala 1475 1485	tct	Ser	
gac aac gcc ttt gtt aaa gga ctt Asp Asn Ala Phe Val Lys Gly Leu 1475	gct	Ala	485
gac aac gcc ttt gtt aaa gga ctt Asp Asn Ala Phe Val Lys Gly Leu 1475	gcc	Ala	
gac aac gcc ttt gtt aaa gga ctt Asp Asn Ala Phe Val Lys Gly Leu 1475	agg	Lys	
gac aac gcc ttt gtt aaa gga ctt Asp Asn Ala Phe Val Lys Gly Leu 1475	gaa	Glu	
gac aac gcc ttt gtt aag Asp Asn Ala Phe Val Lys 1475	H	딦	
gac aac gcc ttt gtt aag Asp Asn Ala Phe Val Lys 1475	gga	Gly	480
gac aac gcc ttt gtt Asp Asn Ala Phe Val 1475	aaa	Š	()
gac aac gcc ttt Asp Asn Ala Phe 1475			
gac aac Asp Asn 1475	ttt	ζŢ	
gac aac Asp Asn 1475	gac	Ala	
gac Asp	aac	Asn	4.
	gac	Asp	 1

4512		
gcc	Ala	
gtt	Val Ala	
gcc	Ala	
aat	Asn	
gat tta aat gcc gtt g	Asp Leu Asn Ala	1500
gat	Asp	` '
ggt	. Gly 1	
a act gtg	ਯ	
act	a Val Thr V	
gta	Val	1495
gca	Ala	` '
gcc	Ala	
aac	rs Asn Ala Ala Val Th	
aga i	Lys 7	
acc	Thr	490
agg	Lys	,—

a Lys Lys	1520
Asp Thr Gly Thr Thr	1515
Thr Phe Ala Gly	1510
Gln Thr Pro	1505
	Thr Pro Leu Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys

4608		
aca gac acc aat aag	Asp Thr Asn Lys	1535
t ttg acc atc aaa ggt ggg caa	Lys Gly Gly Gln	1525



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cta acc gat aat aac atc ggt gtg gta gca ggt act gat ggc ttc act Leu Thr Asp Asn Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr 1540

4704		
aat ctt aac agc gtt aat gca ggt	sn Leu Asn Ser Val Asn Ala	1565
gac cta acc	Ala Lys Asp Leu Thr A	1560
gtc and ctt o	Val Lys Leu A	1555

4752		
s gca aac ggt	Ala	
tet ttt gta gad	Phe Val Asp	1580
atc	Gly Ile Ser	
ගුපය යයය	Glu Lys	1575
aaa att gat	g	
90 CC	Gly Thr L	1570

ASII IIII 1590		3 2 2 3
	1 1 1	

4848		
aaa gat acc	Lys	1615
ytg ggc aaa	Val Gly Lys Gly	1610
ggc aag gtc atc agt aat	Ile Ser Asn	1605

4896		
s aac ttg ttg ggt ctt	Leu Asn Glu Val Arg Asn Leu Leu Gly Leu	1630
itta aac gaa gta cgc	Leu Asn Glu Val Arg	1625
gct gcc aat gta caa cag 'tta	Asn Val Gln Gln	1620



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5184

cag acc aac tat gtg acc

act tat aac gcc gca ggt

gcc ctg ctc gcc Ala Leu Leu Ala

1710

1705

1700

Thr Tyr Asn Ala Ala Gly Gln Thr Asn Tyr Val Thr 1720

4944 4992 5040 5088 5136 Gly Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala Asp Ile Lys 1635 ggt aat gat aac gct gac ggc aat cag gta aac att gcc gac atc aaa aaa gca Lys Asp Pro Asn Ser Gly Ser Ser Asn Arg Thr Val Ile Lys Ala 1650 Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Āsp Thr Glu Lys Leu Āla 1665 act ggt ggt gta caa gtg ggc gtg gat aaa gac ggc aac gct aac ggc Thr Gly Gly Val Gln Val Gly Val Asp Lys Asp Gly Asn Ala Asn Gly 1695 ggc acg gta ctt ggc ggt aaa ggt aat aac gat acc gaa aaa ctt gcc Asp Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly Ser Lys Lys gat tta agc aat gtt tgg gtc aaa acc caa aaa gat ggc agc aaa aaa cca aat tca ggt tca tca tct aac cgc act gtc atc aaa gac



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ttc cat gtc aac ttc cat gtc aac ggc att gac tca ggc att gac tca Gly Ile Asp Ser 1765 cag gcc aag gca Gln Ala Lys Ala 1780 caa gca ggc aac Gln Ala Gly Asn 1795	1735 1740 gat ggc aat caa gag cct gtg gta caa ggg cgt 5280 Asp Gly Asn Gln Glu Pro Val Val Gln Gly Arg 1750 1755	agt gcc tca ggc aag cac tca gtg gcg ata ggt 5328 Ser Ala Ser Gly Lys His Ser Val Ala Ile Gly 1770	gat ggt gaa gcc gcc gtt gcc ata ggc aga caa 5376 Asp Gly Glu Ala Ala Val Ala Ile Gly Arg Gln 1785	caa tcc atc gcc atc ggt gat aac gca caa gcc 5424 Gln Ser Ile Ala Ile Gly Asp Asn Ala Gln Ala 1800	gat caa tcc atc gcc atc ggt aca ggc aat gtg gta gca ggt 5472 Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val Val Ala Gly
ttc cat gtc aac gat ggc aat caa gag cct gtg Phe His Val Asn Asp Gly Asn Gln Glu Pro Val 1750 ggc att gac tca agt gcc tca ggc aag cac tca Gly Ile Asp Ser Ser Ala Ser Gly Iys His Ser 1765 cag gcc aag gca gat ggt gaa gcc gcc gtt gcc Gln Ala Iys Ala Asp Gly Glu Ala Ala Val Ala 1780 caa gca aac caa tcc atc gcc atc ggt gat Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp 1795 1705 ggc gat caa tcc atc gcc atc ggt aca ggc aat Gly Asp Gly Ash Gly Ser Ile Ala Ile Gly Asp Gly Asp Gly Ser Ile Ala Ile Gly Asp	cct gtg (Pro Val 7 755	cac tca (tis Ser '	gtt gcc a	ggt gat a 31y Asp 7	ggc aat g 31y Asn ¹
ttc cat gtc aac gat ggc aat caa g Phe His Val Asn Asp Gly Asn Gln G 1750 ggc att gac tca agt gcc tca ggc a Gly Ile Asp Ser Ser Ala Ser Gly I 1765 cag gcc aag gca gat ggt gaa gcc g Gln Ala Lys Ala Asp Gly Glu Ala A 1780 caa gca ggc aac caa tcc atc gcc a Gln Ala Gly Asn Gln Ser Ile Ala I 1795 ggc gat caa tcc atc gct a Glv Asp Gly Glu Ser Ile Ala I 1795	caa gag (31n Glu]	ggc aag (31y Lys 1 1770	gcc gcc g Ala Ala 7 785	gcc atc g Ala Ile (ggt aca g 31y Thr (
ttc ttc cat gtc aac gat g Phe Phe His Val Asn Asp G 1745 aac ggc att gac tca agt g Asn Gly Ile Asp Ser Ser A 1765 ttc cag gcc aag gca gat g Phe Gln Ala Lys Ala Asp G Thr Gln Ala Gly Asn Gln S acc caa gca ggc aac caa t Thr Gln Ala Gly Asn Gln S acg ggc gat caa tcc atc g	35 gc aat o	rcc tca g la Ser (igt gaa gily Glu I	cc atc ger 11e A	rc atc g la Ile (
ttc cat gtc a bhe His Val A ggc att gac t Gly Ile Asp S cag gcc aag g Gln Ala Lys A 1780 caa gca ggc a Gln Ala Gly A 1795 Glv Asp Gln S	17 ac gat g sn Asp G 1750	ca agt g er Ser A 65	ca gat g la Asp G	ac caa t sn Gln S	cc atc ger er Ile A
ttc ce Phe Hi ggc at Gly Il Gly Il Glh Al Glh Al ggc ga	gtc Val	it gac ti e Asp Si 17	ic aag g a Lys A 1780	ggc Gly	it caa t ip Gln s
i jaga jaga jaga jaga jaga jaga jaga jag		aac ggc at Asn Gly Il	ttc cag gc Phe Gln Al	acc caa gc Thr Gln Al 179	acg ggc ga Thr Gly As

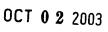


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5520		
aag gct gat aac	Lys Ala	1840
cca agc	Pro Ser Thr	1835
ggt gcc atc ggc	G	1830
aag cac tct	Lys His Ser	1825

0000	5616
agt tac agt gtg ggt aat aac aac cag ttt acc gat gcc act caa acc Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala Thr Gln Thr 1855	gat gtc ttt ggt gtg ggc aat aac atc acc gtg acc gaa agt aac tcg Asp Val Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu Ser Asn Ser 1860
	•

5712		
acc aca gca	Thr	
aca acc	Thr Thr	0061
	Thr Ala Gly	\ 1
gac ggc	Ser Asp Gly Ti	1895
वववं ववव	Lys Lys	
aca caa gcc	Thr Gln Ala	1890
	_	



TECH CENTER 1600/2900

SEP 2 6 2003 SEP 2

2808	5856	5904	5952	0009	6048
gcg gtc tcc gtg ggt gcc tca ggt gct gaa cgc cgt atc caa aat gtg	gca gca ggt gag gtc agt gcc acc agc acc gat gcg gtc aat ggt agc	cag ttg tac aaa gcc acc caa agc att gcc aac gca acc aat gag ctt	gac cat cgt atc cac caa aac gaa aat aaa gcc aat gca ggg att tca	tca gcg atg gcg tcc atg cca caa gcc tac att cct ggc aga	tcc atg gtt acc ggg ggt att gcc acc cac aac ggt caa ggt gcg gtg
Ala Val Ser Val Gly Ala Ser Gly Ala Glu Arg Arg Ile Gln Asn Val	Ala Ala Gly Glu Val Ser Ala Thr Ser Thr Asp Ala Val Asn Gly Ser	Gln Leu Tyr Lys Ala Thr Gln Ser Ile Ala Asn Ala Thr Asn Glu Leu	Asp His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala Gly Ile Ser	Ser Ala Met Ala Ser Met Pro Gln Ala Tyr Ile Pro Gly Arg	Ser Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln Gly Ala Val
1935	1940	1955	1970	1985	2015



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9609 gga ctg tcg aag ctg tcg gat aat ggt caa tgg gta ttt aaa Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp Val Phe Lys 2020 2030 gtg Val gca Ala

ggt Gly

a ggg gcg gca gtt g . Gly Ala Ala Val G 2045 atc aat ggt tca gcc gat acc caa ggc cat gta Ile Asn Gly Ser Ala Asp Thr Gln Gly His Val 2035

gca ggt ttt cac ttt Ala Gly Phe His Phe 2050

6159



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TECH CENTER 1600/2900

Moraxella catarrhalis les1 200kDa

48	96	144	192	240	288
a ggc aca ttt 7 Gly Thr Phe 15	g agt agc agt 7 Ser Ser Ser 30	ctg act cgt y Leu Thr Arg 5	aat ggc agt 1 Asn Gly Ser	aca ggc aac Thr Gly Asn 80	att ggt agt Ile Gly Ser 95
aaa gcc aca Lys Ala Thr	agc gga ggg Ser Gly Gly	gtc atc cgc Val Ile Arg 45	gcg acg ctc Ala Thr Leu 60	ttt ggt acc Phe Gly Thr 75	tcc att gct Ser Ile Ala
ttt aac Phe Asn 10	tcc cac Ser His 25	tct cct Ser Pro	atc ggt Ile Gly	atc gca Ile Ala	gaa gca Glu Ala 90
aaa gtc atc Lys Val Ile	tgc gcc aaa Cys Ala Iys	gtg ggc agc Val Gly Ser 40	atc cta gtg Ile Leu Val 55	aat agc aag Asn Ser Lys 70	gct agc aat Ala Ser Asn
atc tat Ile Tyr 5	gca gag Ala Glu 20	gga cag Gly Gln	ctc gct Leu Ala	caa aat Gln Asn	gcc tcg Ala Ser 85
ATG aat cac Met Asn His 1	atg gcc gtg Met Ala Val	agt acc gca Ser Thr Ala 35	gtt gcc acg Val Ala Thr 50	gct tat gct Ala Tyr Ala 65	aat gac aat Asn Asp Asn



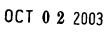
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TECH CENTER 1600/290

SEP 2 6 2003 SEP 2

336	384	432	480	528	576
ctt gct aag gca cat gcc aat caa gct att gct atc ggt ggt agc aaa	cca gat cct cgt aat caa gcg gct aat cag aag gca ggt tcc cac gcc	aaa ggt aaa gag tcc atc gcc atc ggt ggt gat gta ctg gct gag ggt	gat gcc tcg att gcc att ggt agt gat gac tta tat ttg gat agg aat	agc act aac tct aaa tat cca aat ggt ctt ctt agc act ctt att caa	aac cat aca gta tta cgc caa ata cga gac tca aat ggt tct cag aaa
Leu Ala Lys Ala His Ala Asn Gln Ala Ile Ala Ile Gly Gly Ser Lys	Pro Asp Pro Arg Asn Gln Ala Ala Asn Gln Lys Ala Gly Ser His Ala	Lys Gly Lys Glu Ser Ile Ala Ile Gly Gly Asp Val Leu Ala Glu Gly	Asp Ala Ser Ile Ala Ile Gly Ser Asp Asp Leu Tyr Leu Asp Arg Asn	Ser Thr Asn Ser Lys Tyr Pro Asn Gly Leu Leu Ser Thr Leu Ile Gln	Asn His Thr Val Leu Arg Gln Ile Arg Asp Ser Asn Gly Ser Gln Lys
100 110	120	130	145	165	180





TECH CENTER 1600/2900

OFENT & TRACE, RES

624	672	720	768	816	864
tat aga cgc aca gca gca gaa gga cac gcc agt act gca gtg gga gcc Tyr Arg Arg Thr Ala Ala Glu Gly His Ala Ser Thr Ala Val Gly Ala 200	atg gca tat gca aag ggt cat ttt gcc aac gcc ttt ggt aca cgg tca Met Ala Tyr Ala Lys Gly His Phe Ala Asn Ala Phe Gly Thr Arg Ser 210	aca gct gaa ggc aac tat tcc ttg gca gta ggt ctt acc gcc aaa gcc Thr Ala Glu Gly Asn Tyr Ser Leu Ala Val Gly Leu Thr Ala Lys Ala 225	gaa aaa gga tat aca atc gct att ggt tct aat gca caa gct atc aat Glu Lys Gly Tyr Thr Ile Ala Ile Gly Ser Asn Ala Gln Ala Ile Asn 255	tat gga gca cta gcc ctt ggt gca gat act cga gtt gat ttg gat tac Tyr Gly Ala Leu Ala Leu Gly Ala Asp Thr Arg Val Asp Leu Asp Tyr 260	ggt att gcc cta ggt tat ggt tct cag atc ctt aat aat aat aat aat aat Gly Ile Ala Leu Gly Tyr Gly Ser Gln Ile Leu Asn Asn Asn Asn 275
ĬĠ ĬĠ	rd E	B FI CI	D Q .	th FI	OI O



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				•	
912	096	1008	1056	1104	1152
: aat aaa gcc tat gta cca gaa ggt aat ggg tca aac ata aaa 1 Asn Lys Ala Tyr Val Pro Glu Gly Asn Gly Ser Asn Ile Lys 300	aaa gcc acc ggc aat ggt tta ttt tcc att ggt agt agc act Lys Ala Thr Gly Asn Gly Leu Phe Ser Ile Gly Ser Ser Thr 310	g cgt aaa atc atc aat gtc ggt gca ggt tat gag gat acc gat Harg Lys Ile Ile Asn Val Gly Ala Gly Tyr Glu Asp Thr Asp 335	aat gtg gca cag cta aaa gcg gtg gag aat ctg gct aag cgt Asn Val Ala Gln Leu Lys Ala Val Glu Asn Leu Ala Lys Arg 340	act ttt aag ggt gat gat aac ggt act ggc gtt aag aaa aaa Thr Phe Lys Gly Asp Asp Asn Gly Thr Gly Val Lys Lys 355 365	: gag act tta acc att aaa ggt ggt gag acc caa gcg gac aag 7 Glu Thr Leu Thr Ile Lys Gly Gly Glu Thr Gln Ala Asp Lys 375
aat aat Asn Asn 290	tog tot Ser Ser 305	atc aag Ile Lys	gcg gtc Ala Val	caa att Gln Ile	ctg ggc Leu Gly 370
₩ W	3 % tt	at []	N R	8 5	せど



OCT 0 2 2003 TECH CENTER 1600/2900

1200	1248	1296	1344	1392	1440
cta acc gat aat aac att ggt gtg gta aca gat aat aat ggt	ctg aaa gtt aaa ctt gct aaa aac cta agc ggt ctt gaa aca gtt agc	acc aga aga cta acc gcc agc gag aga gtt acg gta ggt agt ggt aat	aac acc gct gag cta caa agc ggt ggt tta acc ttt acc cca aca aca	aat gca agc aca gac aaa acc gtc tat ggc act gat ggg ctt aag ttt	act gat aat tct aat acg gca ctt gaa gat act act cgt atc aca aaa
Leú Thr Asp Asn Asn Ile Gly Val Val Thr Asp Asn Asn Thr Gly	Leu lys Val lys Leu Ala lys Asn Leu Ser Gly Leu Glu Thr Val Ser	Thr Lys Asn Leu Thr Ala Ser Glu Lys Val Thr Val Gly Ser Gly Asn	Asn Thr Ala Glu Leu Gln Ser Gly Gly Leu Thr Phe Thr Pro Thr Thr	Asn Ala Ser Thr Asp Lys Thr Val Tyr Gly Thr Asp Gly Leu Lys Phe	Thr Asp Asn Ser Asn Thr Ala Leu Glu Asp Thr Thr Arg Ile Thr Lys
385	410	420	445	450	465



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1488	1536	1584	1632	1680	1728
gat aaa att ggt ttt agc aat aaa gct ggt aca gtt gat gaa aac aaa	cct tat ctt gat aaa gac aag cta aaa gtt ggc aac agc acc cta aac	aac ggt ggc ttg act gtt aat aac acc att ggt ggt agc aat aaa caa	atc caa gtc ggt gct gat ggc att aaa ttt gcc gat gtg aat gtt aat	gta tca aat gcc gca aaa ttc ggc act act cgt att acc gaa gag gaa	att ggc ttt gct gat gct gat ggt aaa gtt gat aaa aag tca cca tat
Asp Lys Ile Gly Phe Ser Asn Lys Ala Gly Thr Val Asp Glu Asn Lys	Pro Tyr Leu Asp Lys Asp Lys Leu Lys Val Gly Asn Ser Thr Leu Asn	Asn Gly Gly Leu Thr Val Asn Asn Thr Ile Gly Gly Ser Asn Lys Gln	Ile Gln Val Gly Ala Asp Gly Ile Lys Phe Ala Asp Val Asn Val Asn	Val Ser Asn Ala Ala Lys Phe Gly Thr Thr Arg Ile Thr Glu Glu Glu	Ile Gly Phe Ala Asp Ala Asp Gly Lys Val Asp Lys Lys Ser Pro Tyr
495	500	515	530	545 555	565

FIG.5



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1824 gac Asp gca Ala aaa Liys gat Asp acc Thr 590 aaa Liys att Ile gtt Val 605 aat Asn aaa Lys agt Ser gtt Val ggt Gly atc Ile aag Lys ggt Gly 585 caa Gln 600 gtg Val caa Gln gat Asp caa ctt (Gln Leu (ggt Gly gca Ala aat Asn aaa Lys 580 aaa Lys att Ile 595 gat Asp ggc Gly ttg

agt Ser

10/07	192(
rett aga eaa gre eaa	att cgt gat gaa aaa
1 Leu Lys Gln Val Gln	Ile Arg Asp Glu Lys
620	640
aaa cag c	ttc tct a
Lys Gln L	Phe Ser I
6	. 635
act tat	caa agc
Thr Tyr	Gln Ser
gat gca gtc	ggt gcc cta
Asp Ala Val	Gly Ala Leu
615	630
gac gat acc gass Asp Asp Thr 1610	gac gcc gac (Asp Ala Asp (
acg g	caa g
Thr ?	Gln A

1968	2016
ggt cag gaa ttt acg att agt aac ttg tat tct aat ggt aat acc cca Gly Gln Glu Phe Thr Ile Ser Asn Leu Tyr Ser Asn Gly Asn Thr Pro 645 655	מלאם אלם מאס מנה נהי לאי היי ליין ביין

20		
aac ggc atc agt atc	Gly Ile Ser	0.29
acc ttt gag acc atc acc ttt gca	Thr Ile Thr Phe Ala Gly	099



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gac cca : Asp Pro] ggt, gtt Val aaa Lys gtc aaa Lys 680 aaa ggt a Lys Gly 1 gcc gac Asp 675 agc aat g Ser Asn i

7777	2160	2208
agc gat aaa gat ggt	ggt aac gac acc aaa	yc att acc aat gca
Ser Asp Lys Asp Gly	Gly Asn Asp Thr Lys	er Ile Thr Asn Ala
700	720	735
cct aag ctg acc gtg ggt a <u>c</u>	att gag caa gtg gct agc gg	ttg tcc cca aca ctg cct agc
Pro Lys Leu Thr Val Gly Se	Ile Glu Gln Val Ala Ser Gl	Leu Ser Pro Thr Leu Pro Ser
695	710	730
aat ggt ctc acc acg c	aaa act caa ttg gtt a	aac atc att aga gga t
Asn Gly Leu Thr Thr P	Lys Thr Gln Leu Val I	Asn Ile Ile Arg Gly L
690	705	725

2304
gcc gcc agt atc ggt gat ata tta aat aca ggc ttt Ala Ala Ser Ile Gly Asp Ile Leu Asn Thr Gly Phe 760
gac aaa too aaa g Asp Lys Ser Lys A 755



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2352 tcc act tat aac Ser Thr Tyr Asn ttt gtc † Phe Val ; 780 ggc Gly aac cta aaa aat aat agc aac tcc gtt Asn Leu Lys Asn Asn Ser Asn Ser Val 770

2448
aat gtg gat
atc 8
gat
tat
act
ota
agg
aot
acc
Caa
aat
craa acc
gat

2448	
: tat gat gtc aat gtg gat : Tyr Asp Val Asn Val Asp	815
gat Asp	
tat Tyr	
act Thr	810
gta act Val Thr	
aac Lys	-
agt Ser	
acc Thr	
caa	802
aat Asn	
acc Thr	
gaa	
gat	
•	

2496		
aca aac aaa	Gly Lys Thr Asn Lys Ile	830
ggc gat aat	Thr Gly Asp Asn Gly	825
gaa ctc	Glu Lys Thr Ile Glu Leu	. 820

2544		
aat ggt aaa	Asn	845
aat gct	Asn Ala	
aga	Thr Thr	
acc	Thr	840
aca ctg	Thr Leu	
acc acc		÷
agg	Lys	835
ggc gtc	Gly Val I	

la Leu Val Asn Ala Lys Asp	860
Thr Thr Asp Asn Asp	855
Thr Asn Phe Ser	850
	Asn Phe Ser Thr Thr Asp Asn Asp Ala Leu Val Asn Ala Lys



OCT 0 2 2003 TECH CENTER 1600/2900

gcc gaa aat cta aac acc cta gcc aag gan Glu Asn Leu Asn Thr Leu Ala Lys 870 aca gca gac acc gcc cta caa acc ttt 885 act gat gac gaa acc atc acc gtg ggt Thr Asp Asp Glu Thr Ile Thr Val Gly 905 aag acc gtc aac act cta aaa ctc aaa ctc aaa lys Thr Val Asn Thr Leu Lys Leu Lys Ieu Lys Gly Thr Asn Lys Asp Gly Thr Val Thr 930 gct acc aat aaa gat ggt acg gtt acc Ala Thr Asn Lys Asp Gly Thr Val Thr 930 ggt ctt aaa gcc ggc gac agc acc act Gly Irr Thr 750 ggt ctt aaa gcc ggc gac agc acc act Gly Asp Ser Thr Thr 950	2640	2688	2736	2784	2832	2880
	gcc gaa aat cta aac acc cta gcc aag gaa att cac acc acc	aca gca gac acc gcc cta caa acc ttt aaa gtc aaa aaa gac	act gat gac gaa acc atc acc gtg ggt aaa gat ggt aca caa	aag acc gtc aac act cta aaa ctc aaa ggt gaa aac ggt cta	gct acc aat aaa gat ggt acg gtt acc ttt ggc att aac acc	ctt aaa gcc ggc gac agc acc act cta aac aaa gat ggc
	Ala Glu Asn Leu Asn Thr Leu Ala Lys Glu Ile His Thr Thr	Thr Ala Asp Thr Ala Leu Gln Thr Phe Lys Val Lys Lys Asp	Thr Asp Asp Glu Thr Ile Thr Val Gly Lys Asp Gly Thr Gln	Lys Thr Val Asn Thr Leu Lys Leu Lys Gly Glu Asn Gly Leu	Ala Thr Asn Lys Asp Gly Thr Val Thr Phe Gly Ile Asn Thr	Leu Lys Ala Gly Asp Ser Thr Thr Leu Asn Lys Asp Gly
	870	895	900	920	930	950



FIG.5K

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3168

Ile Thr Asn Ile Gln Ser Gly Asp Ile Thr Gln Asn Ser Asn 1045

Lys

aga agg att acc agc att caa tca ggt gat att acc caa aac agc aat

3024 3072 3120 act ggc att Thr Gly Ile 990 ggt gaa gtt gaa att acc aac act ggc att aac gca ggt ggt Gly Glu Val Glu Ile Thr Asn Thr Gly Ile Asn Ala Gly Gly 1030 1030 tct att aaa aac ccc gct agt aac gaa caa atc caa gtc ggt gct gat Ser Ile Lys Asn Pro Ala Ser Asn Glu Gln Ile Gln Val Gly Ala Asp 975 aca agc cgt atc acc aaa gat caa att ggc ttt act ggg gct Thr Ser Arg Ile Thr Lys Asp Gln Ile Gly Phe Thr Gly Ala 995 Asn Gly Ser Leu Asp Thr Thr Lys Pro His Leu Thr Lys Asp Lys Leu 1010 aat ggc tca ctt gat acc acc aaa ccc cac cta acc aaa gac aag ctt ggt aat toa ago Gly Asn Ser Ser ttt gcc aag gtt gat aag Phe Ala Lys Val Asp Lys 980 Gly Val Lys ggc gtg aag ggc g aaa gtg Lys Val 1025 gat Asp



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3216 3408 3456 3264 3312 3360 gat gct gtg aca ggc ggt cgg gtt tat gat tta aaa acc gaa ctt gaa Asp Ala Val Thr Gly Gly Arg Val Tyr Asp Leu Lys Thr Glu Leu Glu 1060 1065 ttc tca gta gca gat gaa caa ggt aat cac ttt acg gtt agt aac cct Phe Ser Val Ala Asp Glu Gln Gly Asn His Phe Thr Val Ser Asn Pro 1090 1095 agc aga atc agc agt gct gct aga aca gca caa agc tca tta cac gaa Ser Lys Ile Asn Ser Ala Ala Lys Thr Ala Gln Asn Ser Leu His Glu 1075 Tyr Ser Ser Tyr Asp Thr Ser Lys Thr Ser Asp Val Ile Thr Phe Ala 1105 ggt gaa aac ggc att acc acc aag gta aat aaa ggt gtg gtg cgt gtg Gly Glu Asn Gly Ile Thr Thr Lys Val Asn Lys Gly Val Val Arg Val 1125 ggc att gac caa acc aaa ggc tta acc acg cct aag ctg acc gtg ggt Gly Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr Val Gly 1140 1150 tac tcc agt tat gac acc tca aag acc tct gat gtc atc acc ttt gca



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aat aat ggc aaa ggc att gtc att gac agt aaa gat ggt caa aat Asn Asn Asn Gly Lys Gly Ile Val Ile Asp Ser Lys Asp Gly Gln Asn 1155 1160

3552		
	n Asp Gly	
acc aat	Thr Asn	
aat gtt	Asn Val	1180
cta gct	Leu Ala	
aac act	Thr	.175
cta agc	Ŕ	 1
aca gga	Thr Gly	
acc atc ;	Thr Ile !	1170
.0		

3600		
gac aaa acc	Asp Lys Thr	1200
lac acc	Sp Thr	
cc aat	la Asn	1195
igg ctt		
Ğ	딩	190
gca cta agc		11
cac		
	Ala G	1185

aac Asn	1215
gat gtg cta aac gca ggc ttt a Asp Val Leu Asn Ala Gly Phe A	1210
ggt	1205

3698		
act tat gac act gtt gac	r Asp Thr Val Asp	1230
ttt gtc tcc	Phe Val Ser	1225
gtt gad	Gly Asn Gly Glu Ala Val Asp	1220

3744		·
ytg acc tat gat	/al Thr Tyr Asp Asp	1245
acc acc gct	Thr Thr	1240
ttt atc gat ggc aat	Phe Ile Asp Gly Asn Ala	1235



FIG.5N

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3792 3840 3888 3936 3984 4032 acc agt aaa gtg gtc tat gat gtc aat gtg gat aat aaa acc Thr Ser Lys Val Val Tyr Asp Val Asn Val Asp Asn Lys Thr att gaa gtg aca agt gat aaa aaa ctt ggc gtc aaa acc acc aca ctg Ile Glu Val Thr Ser Asp Lys Lys Leu Gly Val Lys Thr Thr Thr Leu 1280 acc aga aca agt gct agt ggt agt gca acc aga ttt agt gcc gcc ggt Thr Lys Thr Ser Ala Asn Gly Asn Ala Thr Lys Phe Ser Ala Ala Asp ggc gat gcc ctt gtt aaa gcc agt gat atc gcc acc cat cta aat acc Gly Asp Ala Leu Val Lys Ala Ser Asp Ile Ala Thr His Leu Asn Thr 1300 tat gtg gat gct gat ggc aac aag gtc atc tat gac agt Tyr Val Asp Ala Asp Gly Asn Lys Val Ile Tyr Asp Ser 1295 1260 1340 1275 1290 1255 1270 1285 agc Seg Ser Lys 7 1250 tca gca Ser Ala agc aga



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4080 Gly Gln Val Asp Lys 1360 ggt caa gtg gac aaa Thr Asp Lys Lys Tyr Tyr Gln Val Asn Asp Lys 1345 acc gat aag aag tac tat caa gtc aat gac aag

FIG.50

4128 Asn Lys Glu Val Ala Lys Asp Lys Leu Val Ála Gln Ála Gln Thr Pro 1365 1375 gaa gtt gcc aaa gac aaa ctg gtc gcc caa gcc caa acc cca aac aaa

4176 Gly Thr Leu Ala Gln Met Asn Val Lys Ser Val Ile Asn Lys Glu 1380 ggc aca ttg gct caa atg aat gtc aaa tca gtc att aac aaa gag Asp gat

4224 Gln Val Asn Asp Ala Asn Lys Lys Gln Gly Ile Asn Glu Asp Asn Ala 1395 gta aat gat gcc aat aaa aag caa ggc atc aat gaa gac aac gcc Caa

ttt atc aaa ggg ctt gaa aac gcc gcc aaa gac acc aaa acc aaa aac Phe Ile Lys Gly Leu Glu Asn Ala Ala Lys Asp Thr Lys Thr Lys Asn 1410 gcc gca gta act gtg ggt gat tta aat gcc gtt gcc caa aca ccg ctg Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala Gln Thr Pro Leu 1430 1425



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ggc aca acg gct aaa aaa ctg ggc gag act Gly Thr Thr Ala Lys Lys Leu Gly Glu Thr 1450 s ttt gca ggg gat aca g r Phe Ala Gly Asp Thr G 1445 acc Thr

4416		
aag cta acc gat aat	Lys Leu	1470
caa aca gac acc aat	31n Thr Asp	1465
ttg acc atc aaa ggt ggg o	Leu Thr Ile Lys Gly Gly	1460

1485
1480
1475

4512		
gc acc	Gly Gly Thr Arg	1500
agc gtt aat	Ser Val Asn	
, acc aat ctt aac	hr Asn Leu Asn	1495
aaa gac cta a	Leu	1490

4560		
gcc aaa gca	Ala Lys Ala	1520
ggt caa	Gly Gln	
a gac gca aac	Ala Asr	1515
atc tct ttt gta	Ile Ser Phe Val	1510
gat gaa aaa ggc	Asp Glu Lys Gly	1505

4608		
CGC	Arg	
aaa	Lys	1535
ggc	GLy	₹
ggt	Gly	
ctg	Leu	
gac	Asp	
ctg	Leu	1530
999	G1y	
aat	Asn	
ggc	Ala	
agt	Ser	
cta	Leu	1525
gtg	Val	·
cct	Pro	
aac acc cct gtg cta agt gcc aat ggg ctg gac ctg ggt ggc aaa cgc	Thr	
aac	Asn	



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atc agt aac atc ggt gca gct gtt gat gat aac gat gcg gtg aac ttt Ile Ser Asn Ile Gly Ala Ala Val Asp Asp Asn Asp Ala Val Asn Phe 1540 1550

4704		
iac aac cta aac aac caa	: Thr Val Asn Asn Leu Asn Asn Gln	1565
aac	1,7	1560
aag cag ttt aat gaa gtt gcc		1555

4752		
cc aat	e Val Val Thr Asp Ala Asn Gly	1580
a ccc ttt g	Phe V	
gog toa tti	Ala Ser Lei	1575
agt aac tca ggt	Ser Asn Ser Gly ,	1570

4800		
aag	lle Lys Gly	1600
aag ccc caa aaa gcc	Lys Pro Gln Lys Ala	1595
ic acc gat ggc	Thr Asp Gly	1590
	Lys Pro Ile	1585

4848		
cct gtg gac	Pro Val Asp	1615
ggc gta	Gly Val	
aac gcc aac	Asn Ala Asn	1610
cac dcc	Ala	
aaa tac tat	Gly Lys Tyr Tyr His .	1605
gcc gat ggt	Ala Asp Gly	

4896		
tg ga	eu Ala	
aat c	ns Pro Ile Thr Asp Ala Asp Lys Leu Ala Asn Leu Al	1630
t gac	u Ala	
aa ct	lys Lei	
gaca	Asp Li	
; gag) Ala	1625
Sc gat	or Asp	
atc 'a	lle II	
) (CCC	Pro	
ggc aag	Gly Lys	1620
gat gg	Sp Gl	
aaa ge	Lys As	



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yct cat ggc aaa ccc ctt ga

4944	4992	5040	2088	5136	5184
gct cat ggc aaa ccc ctt gat gca ggt cat caa gtg gtg gca agc cta Ala His Gly Lys Pro Leu Asp Ala Gly His Gln Val Val Ala Ser Leu 1635 1645	ggc ggc aac tca gat gcc atc acc cta acc aac atc aag tcc act ttg Gly Gly Asn Ser Asp Ala Ile Thr Leu Thr Asn Ile Lys Ser Thr Leu 1650	cca caa att gac aca cca aac aca ggt aat gcc aat gca ggg caa gcc Pro Gln Ile Asp Thr Pro Asn Thr Gly Asn Ala Asn Ala Gly Gln Ala 1665 1670 1680	caa agt ctg ccc agc cta tca gca gca cag caa agt aat gct gcc agt Gln Ser Leu Pro Ser Leu Ser Ala Ala Gln Gln Ser Asn Ala Ala Ser 1695	gtc aaa gat gtg cta aat gta ggc ttt aac ttg cag acc aat cac aat Val Lys Asp Val Leu Asn Val Gly Phe Asn Leu Gln Thr Asn His Asn 1700	caa gtg gac ttt gtc aaa gcc tat gat acc gtc aac ttt gtc aat ggt Gln Val Asp Phe Val Lys Ala Tyr Asp Thr Val Asn Phe Val Asn Gly 1715



FIG.5S

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5232 5328 5280 aca ggt gcc gac atc aca agc gtg cgt agt gct gat ggc acg atg agt Thr Gly Ala Asp Ile Thr Ser Val Arg Ser Ala Asp Gly Thr Met Ser 1730 1735 aac atc acc gtc aac acc gcc tta gca gcg acc gat gat gat ggc aat Asn Ile Thr Val Asn Thr Ala Leu Ala Ala Thr Asp Asp Asp Gly Asn 1745 1745 gtg ctt atc aaa gcc aaa gat ggt aag ttc tac aaa gca gac gac ctc Val Leu Ile Lys Ala Lys Asp Gly Lys Phe Tyr Lys Ala Asp Asp Leu 1765

5376

atg cca aac ggc tca cta aaa gca ggc aaa tca gcc agt gat gcc aaa Met Pro Asn Gly Ser Leu Lys Ala Gly Lys Ser Ala Ser Asp Ala Lys 1780 1790

5424 5472 act cca act ggt cta agc ctt gtt aac ccc aat gct ggt aaa ggc agt Thr Pro Thr Gly Leu Ser Leu Val Asn Pro Asn Ala Gly Lys Gly Ser 1795 1805 aca ggc gat gca gtg gct ctt aat aac tta tca aaa gcg gta ttt aaa Thr Gly Asp Ala Val Ala Leu Asn Asn Leu Ser Lys Ala Val Phe Lys 1810



FIG.51

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5616 5568 1840 aat gta ggc ggt aag gtc atc agc aat gtg ggt aaa ggc aca aaa gac Asn Val Gly Gly Lys Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp 1860 1860 Ile Gln Gly Lys Asp Asn Ser Ser Ile Thr Leu Ser Lys Asp Gly Leu 1845 tct gat ggc atc agt Ser Asp Gly Ile Ser atc caa ggc aaa gat aac agc agc atc acc cta agc aaa gat ggg ctg tcc aaa gat ggt aca act act acc aca gta agc Ser Lys Asp Gly Thr Thr Thr Thr Val Ser 1825

	•
5712	5760
gta aac. Val Asn	aac cgc Asn Arg 1920
t cag n Gln	a tct r Ser
gc aat ly Asn 00	tca tca Ser Ser
gac ggc Asp Gly 1900	tca ggt te Ser Gly Se 1915
gct Ala	
c aac o Asn	a aat O Asn
at gat an Asp 35	ac cca op Pro
ggt aat Gly Asn 1895	iaa gac iys Asp 110
gct Ala	aaa aaa g Lys Lys A 1910
aat Asn	atc Ile
t ggt 1 Gly 1	s gac A Asp
ggt ctt g Gly Leu (1890	att gcc Ile Ala 1905
86 5	at 11

5664

acc gac gct gcc aat gta caa cag tta aac gaa gta cgc aac ttg ttg Thr Asp Ala Ala Asn Val Gln Gln Leu Asn Glu Val Arg Asn Leu Leu 1875



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act gtc atc aaa gca ggc acg gta ctt ggc ggt aaa ggt aat aac gat Thr Val Ile Lys Ala Gly Thr Val Leu Gly Gly Lys Gly Asn Asp 1935

2856		
gac	Asp	
agg	Asp Lys	
gat	Asp	1950
gtg	Val	
ğ	Gly	
gtg	Val	
įд	Gln	
gta	/al	1945
ggt	Gly 1	
ggt	GLY	
act	Thr	
ctt gcc	Ala	
ctt	Leu	1940
agg	Lys	(-1
gaa	Thr Glu	
acc	Thr	

5904		
agg	Trp Val Lys Thr Gln Lys	1965
gat tta agc aat gtt	Asp Leu Ser Asn Val Trp Val	1960
ggc aac gct aac ggc	Gly Asn Ala Asn Gly Asp Leu S	1955

5952		
ac gcc gca ggt cag	Ala Ala Gly	30
ctc gcc act tat aac	112	1980
aaa gcc ctg	s Lys Ala Leu Leu Ala	1975
gat ggc agc aaa	Asp Gly Ser Lys	1970

0009		
aga ata aat	Arg Ile Asn	2000
att	. Ile Asp Arg	
ga	a Glu Ala 🛚	1995
c ccc ga	n Pro Ala	
raccaacaac	Tyr Leu Thr Asn Asn	1990
tat ttg	-	I
acc aac	Thr Asn	1985

6048	
aat caa gag cct Asn Gln Glu Pro	
t gtc aac gat ggc s Val Asn Asp Gly	2010
gaa caa ggt atc cgc ttc ttc cat gan Glu Glu Glv Ile Ard Phe Phe His '	2005



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SEP 2 6 TOTAL SEPTEMBER TRADERS

9609	6144	6192	6240	6288	6336
gtg gta caa ggg cgt aac ggc att gac tca agt gcc tca ggc aag cac	tca gtg gcg ata ggt ttc cag gcc aag gca gat ggt gaa gcc gcc gtt	gcc ata ggc aga caa acc caa gca ggc aac caa tcc atc gcc atc ggt	gat aac gca caa gcc acg ggc gat caa tcc atc gcc atc ggt aca ggc	aat gtg gta aca ggt aag cac tct ggt gcc atc ggc gac cca agc act	gtt aag gct gat aac agt tac agt gtg ggt aat aac aac cag ttt atc
Val Val Gln Gly Arg Asn Gly Ile Asp Ser Ser Ala Ser Gly Lys His	Ser Val Ala Ile Gly Phe Gln Ala Lys Ala Asp Gly Glu Ala Ala Val	Ala Ile Gly Arg Gln Thr Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly	Asp Asn Ala Gln Ala Thr Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly	Asn Val Val Thr Gly Lys His Ser Gly Ala Ile Gly Asp Pro Ser Thr	Val Lys Ala Asp Asn Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Ile
2020	2035	2050 2050	2065 2080	2095	2100



TECH CENTER 1600/2900

SEP 2 6 2003 SUE

6384	6432	6480	6528	6576	6624
gat gcc act cag acc gat gtc ttt ggt gtg ggc aat aac atc acc gtg Asp Ala Thr Gln Thr Asp Val Phe Gly Val Gly Asn Asn Ile Thr Val 2115	acc gaa agt aac tog gtt gcc tta ggt toa aac tot gcc atc agt goa Thr Glu Ser Asn Ser Val Ala Leu Gly Ser Asn Ser Ala Ile Ser Ala 2130	ggc aca cac gca ggc aca caa gcc aaa aaa	aca acc aca gca ggt gca aca ggt acg gtt aaa ggc ttt gct gga Thr Thr Thr Ala Gly Ala Thr Gly Thr Val Lys Gly Phe Ala Gly 2175	caa acg gcg gtt ggt gcg gtc tcc gtg ggt gcc tca ggt gct gaa cgc Gln Thr Ala Val Gly Ala Val Ser Val Gly Ala Ser Gly Ala Glu Arg 2180	cgt atc caa aat gtg gca gca ggt gag gtc agt gcc acc agc acc gat Arg Ile Gln Asn Val Ala Ala Gly Glu Val Ser Ala Thr Ser Thr Asp



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	6720	6768	6816	6864	6912
gcg gtc aat ggt agc cag ttg tac aaa gcc acc caa ggc att gcc aac Ala Val Asn Gly Ser Gln Leu Tyr Lys Ala Thr Gln Gly Ile Ala Asn 2210	gca acc aat gag ctt gac cat cgt atc cac caa aac gaa aat aaa gcc Ala Thr Asn Glu Leu Asp His Arg Ile His Gln Asn Glu Asn Lys Ala 2225 2240	aat gca ggg att tca tca gcg atg gcg atg gcg tcc atg cca caa gcc Asn Ala Gly Ile Ser Ser Ala Met Ala Met Ala Ser Met Pro Gln Ala 2255	tac att cct ggc aga tcc atg gtt acc ggg ggt att gcc acc cac aac Tyr lle Pro Gly Arg Ser Met Val Thr Gly Gly Ile Ala Thr His Asn 2260	ggt caa ggt gcg gtg gca gtg gga ctg tcg aag ctg tcg gat aat ggt Gly Gln Gly Ala Val Ala Val Gly Leu Ser Lys Leu Ser Asp Asn Gly 2275	caa tgg gta ttt aaa atc aat ggt tca gcc gat acc caa ggc cat gta Gln Trp Val Phe Lys Ile Asn Gly Ser Ala Asp Thr Gln Gly His Val

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ttt Phe ggg gcg gca gtt ggt gca ggt ttt cac Gly Ala Ala Val Gly Ala Gly Phe His 2305

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OF E JC130 SER 18 TRADERS

Alignment of amino acid sequence of 200kDa proteins of M. catarrhalis strains

FIG.6A

4223 Q8 LES-1	4223 Q8 LES-1	
10 20 30 40 50 60 MNHIYKVIFNKATGTFNAVAEYAKSHSTGGGSCATGQVGSVCTLSFARLAALAVLVIGATLSGS R R R R R R R	110 120 130 140 150 160 ENANAQGQAIAIGSSNKTVNGSSLD-KIGTDATGQESIAIGGDVKASGDASIAIGSDDLHLLD SLSKSVKPDP.NG.NG-NV.SH.K.NL.EY.PK SL.K.HANG.KPDPRNQAANQ.A.SH.K.KL.EY.DR 170 180 200QHGNPKHPKGTLINDLINGHAVIKEIRSSKINDVKYRNLDLNETHKHEIK.QT.T.GKINST.S.Y.N.L.STQN.TRQD.NGSQ	210 220 240 250 260 RITASCHASTAVGAMSYAQGHFSNAFGTRATAKSAYSLAVGLAATAEGQSTIAIGSDATSSSIG R.Q
10 20 MMHIYKVIFNKATGTFMAVAEYAK	110 120 ENANAQGQAIAIGSSNKTVN SLSKSVKPDP. SL.K.HANG.KPDPF	220 RTTASGHASTAVGAMSYAQGHFSN R.QA.KA.



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4223	. 4223	4223
Q8	Q8	Q8
Les-1	LES-1	LES-1
270 280 300 300AIALGAGTRAQLQGSIALGQGSVVTQSDNN-SRPAYTT.I.GN.VVN.GRGVFQILDRTDASVLDVD.DYGYQILNNNNNNKV	310 320 350 FNTQALDPKFQAITNITKAGPL-SIG-SNSIKRKIINVGAGVAKTDAVNVAQLEAVVKW LGKT.ADQYKRQGDSTDIFN.NNNSRSRDKL.EEL EGNGSNIKSS-KGNG.FSSTYEDKENL 360 370 380 390 400 AKERRITFQGD-DN-STDVKIGLDNTLTIKGGAETNALIDNN-IGVV N.KK.GN.NS.ERGD.QEA	410 420 430 440 450 460 KEADNSGLKVKLAKTTANLITEANTTTTANKVGSSSSTTAELLSDSLIFTQPNTGSQSTSK TDGNE.TGSS.NKIT.SNTNNNNNQ.GGS-ITKD TDAN-TN.SG.ET.S.KN.T.SEK.TGNNQ.GGT.NAD 470 480 490 500TVYGANGVKFTNNAETTAALGTTRDKIGFARDGTVYGANGVKFTNNAETTAALGTTRLTRDKIGFARDGTVYGANGVKFTNNAETTAALGTTRLTRDKIGFARDGTDLLDSNSI.TKKKGTN



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SEP 2 6 2003 JULY SEP 2 6 2003

	4223	Q8 LES-1			٠.	4223 Q8	LES-1			
. GVDESKPYLDNEKLKVGNSTLNSGSLTVANTTGAKQIQVGANGIKFATVANNVANTSATVG GTVDENKPYLDKDKLKVGNSTLNNGGLTVANTIGGSNKQIQVGADGIKFADVANVNVSN-AAKFG	510 520 530 530 530 530 530 530 530 530 530 53	TARITEEKIGFAGINDGERR.ET.SN. TIRITEEEIGFADADGKKSQG.KK.SN.	540 550 560 570 580 590	SSANDAVTIEQLKAAKPILNAGAGISVIPIEISVDAKSGNVIAPIY IANTKDDINSNNGDLVDSI.T	009	EINSDG		610 620 630 640 650 660	TSDKFSVKGSGINNSLVTAEHLASYINEVNRTADSALQSFTVKEEDDDDANAITVAKDTTKN GNNSNAHDKDDKEPK.QNGNSNG	SIRDKGQEFT.SNLYSNGNT



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4223 Q8 LES-1	4223 Q8 LES-1	
670 680 700AGAVSIIKIKGKNGLTVATKKD-GTVTFGLSQDSGLTIGGKTFNTEVNIT.NRATID.SNTPPNTFETITFA.EISISNDIAK.K.KV.IDPINTP	710 720 730 740 750 KSTLANDGLIVKDINEQIQVGANGIKFTININGSNRGIGIANTARITRDKIGFA L. VGSDIN NR-LV-I VP-SADG. ST. NI IR	810 820 840 850 860 ADQSS-RNIELGNTIQ-DKDKSNAASINDILNTGFNLKANNANPIDFVSTYDIVDFANGNATTAT SP.GAE-ED.VAGKDKTID



890

880

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Q8 LES-1	4223 Q8 LES-1	422. Q8 LES
$Y \cdot E \cdot Q \cdot \dots A \cdot \dots \cdot EK \cdot E \cdot \dots \cdot G \cdot KQ - \dots $	910 920 930 940 950 960 TKINKTSANGNTATINENVINSSDED-ALVINAKDIAENINTIAKEIHTITKGTADTALQITTVKKVD I.TE.TTT.D.HK.SGE	1010 1020 1030 1040 1050 1060 DKNGTVTFGINTTSGLKAGKST-INDSGLSIKNPTGSEQIQVGADGVKFAKVININGVVGAGIDG



OCT 0 2 2003

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	4223 08 1.ES-1		4223 . Q8 . LES-1
1110 1120 1130 1140 1150 1160 TNIQSGELAQNSHDAVTGGKIYDLKTELENKISSTAKTAQNSLHEFSVADEQGNNFTVSNPYSS K.	D.T. N. RV. S. N.A. H. 1190 1200 1200 1180 1190 1200 YDTSKTSDVITFAGENGITTKVNKGVVRVGIDQTKG	1210 1220 1240 1250 1260 LITPKLIVGNNNSKGIVIDSQNGQNITITGLSNITANVINDKGSVRITTEQGNIIKDEDKIRAASI	



.EUR CENTER 1600/2900

O S TRACE STRACES

1330 1340 1350 1360 DDTTIEVK-DKALGVKTTTTLTSTGTGANKFALSNQATGDALVKAS. NK. TS		4223 Q8 LES-1		4223	LES-1
1310 1320 KVTYDDTSKTSKVVYDVNVI 1410 1420 YDSTDNKYYQAKNDGTVDKT KVNDK.Q	1320 1330 1340 XVYDVNVDDTTIEVK-DKKLGVKTTTLTSTGTGAN		1420 ANDGIVDKIKEVA		INK.T



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4223 Q8 LES-1 4223 Q8 LES-1

...DGKYYHANANGVPVDKDGKPITDADKLANLAAHGKP GGKRI SNIGAAVDDNDAVNFKQFNEVAKIVMNIMQSNSGASLPFVVIDANGKPINGIDGKPQKAIKGA. ... DKGVSFVDSSGQAKANTPVLSANGLDL-1590 1580

LDAGHQVVASLGGNSDALTLTNIKSTLPQIDTPNTGNANAGQAQSLPSLSAAQQSNAASVKDVL.

FIG.6H

FAGDIGITTAKKIGETLITIKGGQIDINKLIDNINIGWAGIDGFIVKLAKDLININSWAGGIKLD

1560

1550

1540

1530

1520

1510



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4223 08 TSVR LES-1		Q8 VSSD LES-1		4223	08 1-25-1
NVGFNLQINHNQVDFVKAYDIVNFVNGIGADITSVR	INITVNITALAATIDDICENVLIKAKDGKEYKADDIMPNGSLKAGKSASDAKTIPTGLSLVN.	PNAGKGSTGDAVALNNLSKAVFKSKDGTTTTTVSSD	1600 GGKVISNVG	DNSSITLSKDGLNV	

FIG.6

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OE JC	130
6	Mr JE
\$22	EMARK
PATENT	& TRADE

	4223 Q8 LES-1		4223 Q8 LES-1
1610 1620 1630 1640 1650 KGTKDTDAANVQQINEVRNLLGLGNAGNDNADGNQVNLADIKKDPNSGSSSNRTVTKAGTVLGG	1670 1680 1690 1700KGNNDTEKLATGGIQVGVDKDGNANGDLSNVMVKTQ .V	1710 1720 1730 1740 1750 1760 KDGSKKALLATYNAAGQINYLINNPAEAIDRINEQGIRFFHVNDGNQEPVVQGRNGIDSSASGK .V	1770 1780 1800 HSVALGFQAKADGEAAVALGRQTQAGNQSIAIGDNA



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4223 Q8 Les-1		4223 08 LES-1
	•	

422 98 LES	422	Q8 LES
1810 1820 1840 1860 QATCDQSIAIGTGNVVAGKHSGAIGDPSTVKADNSYSVGNUNQFTDATQIDVFGVGNUITVTES. T. I870 1880 1890 1900 .NSVALGSNSAISAGTHAGTQAKKSDGTAGTTTTAGA	1910 1920 1940 1950 1960 TGTVKGFAGQTAVGAVSVGASGAERRIQNVAAGEVSATSTDAVNGSQLYKATQSIANATNELDH G. 1970 1980 1990 2000 RTHONENKANAGTSSAMAMASMPOAYIRGRSMVTGG	



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4223 Q8 LES-1

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Construction of Plasmids Expressing Portions of the 200 kDa Protein Gene from Strain 4223

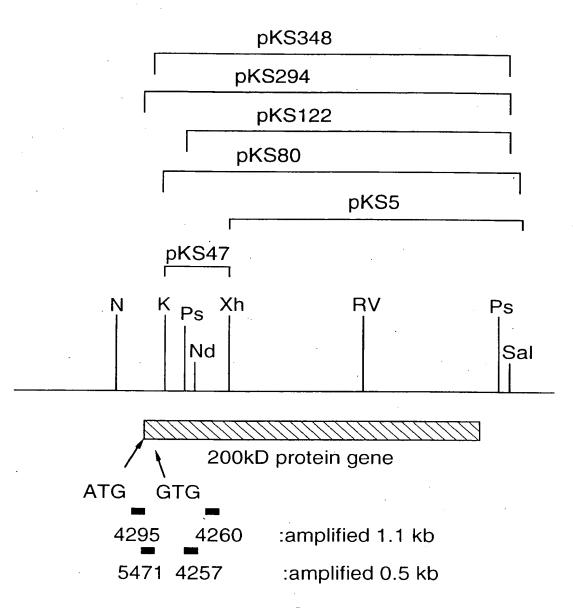


FIG.7



007 0 2 2003

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O FE JC130 SEP 26 MED 13140 NOW

M. catarrhalis M56 200kDa gene in pKS348.

48	96	144	192	240
ATG atc ggt gca acg ctc agt ggc agt gct tat gct caa aaa aaa gat	acc aaa cat atc gca att ggt gaa caa aac cag cca aga cgc tca ggc	act gcc aag gcg gac ggt gat cga gcc att gct att ggt gaa aat gct	aac gca cag ggc ggt caa gcc atc gcc atc ggt agt agt aat aaa act	gtc aat gga agc agt ttg gat aag ata ggt acc gat gct acg ggt caa
Met Ile Gly Ala Thr Leu Ser Gly Ser Ala Tyr Ala Gln Lys Lys Asp	Thr Lys His Ile Ala Ile Gly Glu Gln Asn Gln Pro Arg Arg Ser Gly	Thr Ala Lys Ala Asp Gly Asp Arg Ala Ile Ala Ile Gly Glu Asn Ala	Asn Ala Gln Gly Gly Gln Ala Ile Ala Ile Gly Ser Ser Asn Lys Thr.	Val Asn Gly Ser Ser Leu Asp Lys Ile Gly Thr Asp Ala Thr Gly Gln
1 1 1 1 15 15	20 30	35 46	50 60	65 75 80



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O SE 76 ME TENT & TRACE

288	336
gag tcc atc gcc atc ggt ggt gat gta aag gct agt ggt gat gcc tcg Glu Ser Ile Ala Ile Gly Gly Asp Val Lys Ala Ser Gly Asp Ala Ser 85	att gcc atc ggt agt gat gac tta cat ttg ctt gat cag cat ggt aat Ile Ala Ile Gly Ser Asp Asp Leu His Leu Leu Asp Gln His Gly Asn

384		
ctt att aac		
aac	Ile Asn Asp	
	Leu	120
cct aaa cat ccg aaa ggt a	Lys His Pro Lys Gly	. 115

105

100

432
tat Tyr
aaa Lys '
gta d
gat Asp
aat Asn 140
gat Asp
aag Lys
tca Ser
agc tca Ser Ser
oga Arg 135
ata Ile
gaa Glu
aaa Liys
tta aaa Leu Lys
gta Val 130
gca
·

480		
ggc	31y Ala Met	160
gca gtg	Thr Ala Val	155
gga cac gcc	Ala	20
cgc aca acc gca	Arg Thr Thr Ala	5
aga	Arg	14

528	
ggt aca cgg gca aca Gly Thr Arg Ala Thr	175
aac gcc ttt Asn Ala Phe	170
tca tat gca cag ggt cat ttt tcc Ser Tyr Ala Gln Gly His Phe Ser	165



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576 gag Glu gcc Ala aca Thr 190 gcc r ggt ctt gcc g Gly Leu Ala A gtg (Val (185 gca Ala ttg Se to tat Tyr gcc Ala 180 agt Ser gct aaa a Ala Lys S

624		
og ttg	er Leu	,
age tog t	Ser S	iò
aca tct	Thr Ser	205
gg	Ala	
tct	Ser Asp	
it ggt	e Gly	200
gct att	Ala	
atc	Thr Ile	
tct a	Ser	195
gc caa	Gly Gln	
Q	G	

672		
Cag	G1y	-
gct cag	Ala Gln	220
act cgt	Thr Arg	
ggt gca ggt	ly Ala Gly	215
gcc ctt	Ala Leu	
gga gog ata	Gly Ala Ile	210

720		
aat aat tct	sn Asn Ser	240
agt gat	Asp	235
gtt gtc act	al Thr	
caa ggt tct	Gln Gly Ser	230
	Ile Ala Leu Gly	225
	•	

768		
aag	Pro Lys Phe Gln	255
cag gca cta	Gln 7	250
cca aat	Arg Pro Ala Tyr Thr Pro Asn Thr	245

816		
att ggt agt aac tct	Ile Gly Ser Asn	270
ggt cca ctt tco	Gly Pro Leu Ser	265
acc aat aat acg	Asn Thr Lys	260



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864 aaa acc gat . Lys Thr Asp gtt aat a Val Asn I 285 gca ggt g Ala Gly ggt (gtc Val 280 atc aat o Ile Asn atc Ile aaa Lys ogt Arg 275 atc aaa Ile Lys

FIG.8D

912
y aag tgg gct aag gag 1 Lys Trp Ala Lys Glu 300
gaa gcg gtg gtg Glu Ala Val Val
gca cag cta Ala Gln Leu 295
gcg gtc aat gtg Ala Val Asn Val 290

096		
ata	Ile	320
agg	Lys	
gta		
gac	Asp	
act	Thr	
agt	Ser	315
aac	Asn	
gat	Asp	
gat		
ggt	GLY	
Cag G	_	310
ttt	Phe	
act	Thr	
att	Пе	
	Arg	
gt	Arg	305

1008		
gga		
	Asn	335
acc	Thr	
gag	Glu	
gg	Ala	
ggt		
ggt	GLy	330
	Lys	
att	Ile	
act	Thr	
tta	Leu	
act	먎	325
aat	Asn '	
gat	Asp	
ttg	Leu	
ggt	GLy	

1056		
	Seg	350
gag gct	Val Lys Glu Ala	345.
atc ggt gtg	Asn Ile Gly Val	
tta acc gat aat	Leu Thr Asp Asn	340

1104		
aat ctt act gag	Thr Leu Asn Asn Leu Thr Glu Val Asn	365
gct aaa act tta aac	25.	360
ctg aaa gtt aaa ctt	Leu Lys Val Lys Leu Ala I	355



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FIG.8E

1152	1200	1248	1296	1344	1392
aca act aca tta aat gcc aca acc aca gtt aag gta ggt agt agt agt agt agt ag	agt act aca gct gaa tta ttg agt gat agt tta acc ttt acc cag ccc	aat aca ggc agt caa agc aca agc aaa acc gtc tat ggc gtt aat ggg	gtg aag ttt act aat aat gca gaa aca aca gca atc ggc act act	cgt att acc aga gat aaa att ggc ttt gct cga gat ggt gat gtt gat	gaa aaa caa gca cca tat ttg gat aaa aaa caa ctt aaa gtg ggt agt
	Ser Thr Thr Ala Glu Leu Leu Ser Asp Ser Leu Thr Phe Thr Gln Pro	Asn Thr Gly Ser Gln Ser Thr Ser Lys Thr Val Tyr Gly Val Asn Gly	Val Lys Phe Thr Asn Asn Ala Glu Thr Thr Ala Ala Ile Gly Thr Thr	Arg lle Thr Arg Asp Lys lle Gly Phe Ala Arg Asp Gly Asp Val Asp	Glu Lys Gln Ala Pro Tyr Leu Asp Lys Lys Gln Leu Lys Val Gly Ser
	385 395 400	415	420	445	450



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gca Ala 475 Asp gat Ile att ggc G1yAsn aat gac Asp 470 Ile ata acci Thr gga gtt Val

FIG.8F

1440

Ile

Lys

Lys

Asn

ggt Gly

aag

agg

aat

1488 480 gaa Glu atc Ile 495 acc 캶 gtt Val gog Ala Asp gat Asn 490 aac gct Ala Ser agt agc Ser ggt Gly aaa Lys 485 Ala gcc E ctt aat Asn agt Ser 465

1536 ser Ser atc Ile 995 G1y 510 gct Ala ggc Gly gca Ala Asn agc tta Leu 505 act Ihr Pro cct Lys aag gcc Ala gcc Ala 500 Lys agg Teg Teg ctc Gln Sag

acc Thr

1584

gtt Val aat Asn 9gc G1y 525 Se se aag Lys gct Ala gat Asp gtt Val 520 Ser tg ata Ile gaa Glu act Ihr Pro 515 cct Thr aca gtc Val

1632 gat Asp agt Ser agc Asn ct Lea gag Glu 540 acc Thr acc Thr agg Lys gtg Val 99c G1y 535 Ile aac Asn tac \vec{T} act Thr Pro 530 gcc Ala

agc Asn aat aac Asn acg Thr agg gat agt act

1680 Ser 560 ggt Gly agt Ser 555 ggt Gly aag Lys gtt Val Se set ttt Phe 550 Lys Asp Ser ggc Gly 545



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tat cta aat gaa gtc aat cga Tyr Leu Asn Glu Val Asn Arg 570 575 cat ttg gca agc His Leu Ala Ser tta gtt acc gcc gaa c Leu Val Thr Ala Glu H 565

1776	
: aaa gaa gaa gac gat . Iys Glu Glu Asp Asp	290
caa agc ttt acc gtt Gln Ser Phe Thr Val	585
acg gct gac agt gct cta Thr Ala Asp Ser Ala Leu	

1824	
gat acg aca aaa aat gcc Asp Ihr Ihr Iws Asn Ala	605
acc gtg gct aaa Thr Val Ala Lys	1
gat gac gcc aac gct atc Asp Asp Ala Asn Ala Ile	I

1872
aaa aac ggt cta acg gtt Lys Asn Gly Leu Thr Val 620
aaa ggt Lys Gly
tta aaa ctc a Leu Lys Leu I 615
agc atc t Ser Ile L
ggc gca gtc a Gly Ala Val 610

1920		
gat	Gln Asp Ser	_
ctt	Gly Leu Ser	635
gtt acc ttt	Pr	
ggt acg	Gly Thr	630
aaa gat	Lys Lys Asp	
gct acc a	Thr	625

1968	
igo tto ily Leu	655
aac gat (Asn Asp (
cta aac Leu Asn	650
acc Thr	
ggc aaa agc Gly Lys Ser	345
acc att Thr 116	9
ggt ctg Gly Leu	





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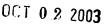
2016	2064	2112	2160	2208	2256
aaa gat acc aac gaa caa atc caa gtc ggt gct aat ggc att aaa ttt Lys Asp Thr Asn Glu Gln Ile Gln Val Gly Ala Asn Gly Ile Lys Phe 660 670	act aat gtg aat ggt agt aat cca ggt act ggc att gca aat acc gct Thr Asn Val Asn Gly Ser Asn Pro Gly Thr Gly Ile Ala Asn Thr Ala 675 685	cgc att acc aga gat aaa att ggc ttt gct ggt tct gat ggt gca gtt Arg Ile Thr Arg Asp Lys Ile Gly Phe Ala Gly Ser Asp Gly Ala Val 690	gat aca aac aaa cct tat ctt gat caa gac aag cta caa gtt ggc aat Asp Thr Asn Lys Pro Tyr Leu Asp Gln Asp Lys Leu Gln Val Gly Asn 705	gtt aag att acc aac act ggc att aac gca ggt ggt aaa gcc atc aca Val Lys Ile Thr Asn Thr Gly Ile Asn Ala Gly Gly Lys Ala Ile Thr 735	ggg ctg tcc cca aca ctg cct agc att gcc gat caa agt agc cgc aac Gly Leu Ser Pro Thr Leu Pro Ser Ile Ala Asp Gln Ser Ser Arg Asn 740

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2304	2352	2400	2448	2496	2544
ata gaa ctg ggc aat aca atc caa gac aaa gac aaa tcc aac gct gcc Ile Glu Leu Gly Asn Thr Ile Gln Asp Lys Asp Lys Ser Asn Ala Ala 755	agc att aat gat ata tta aat aca ggc ttt aac cta aaa aat aat aac Ser Ile Asn Asp Ile Leu Asn Thr Gly Phe Asn Leu Lys Asn Asn Asn 770	aac ccc att gac ttt gtc tcc act tat gac att gtt gac ttt gcc aat Asn Pro Ile Asp Phe Val Ser Thr Tyr Asp Ile Val Asp Phe Ala Asn 785 795	ggc aat gcc acc gcc aca gta acc cat gat acc gct aac aaa acc Gly Asn Ala Thr Ala Thr Val Thr His Asp Thr Ala Asn Lys Thr 805	agt aaa gtg gta tat gat gtg aat gtg gat gat	aca ggc act gat gac aat aaa aaa ctt ggc gtc aaa acc acc aaa ctg Thr Gly Thr Asp Asp Asn Lys Lys Leu Gly Val Lys Thr Thr Lys Leu 835



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2784 2640 2688 2736 2832 2592 aat aat agg Asn aat 880 gac Asp Asn Asn agg Lys Glu Asn Asn aat Asn Ile ggg gga Ala 895 aat ttt aat gtt Val Asn 910 Ala gga Asn Ala agg Thr aat aat ည္တ Leu Asn Пе Glu ggc Asn 925 ctt Phe atc ggc ggg G1YLys Asp Asp ggt Gly 940 ggc Asn Lys gac agg gat aag 860 Asn act Thr Caa Gln agc Thr aaa Lys 875 ಇ೦೦ gta Val Glu Lys acc Ihr ggt Gly gg Ala Ala aag gaa ggc 890 His Lys 905 ggt Gly Asn ပ္တ agg gtg Val gg aac Thr gtt aat Asn att Ile Val Thr 920 agg Lys Val acc gtt ctc Leu 935 Lea atc Ile ctt Glu Thr ggt Gly 855 ggg acc Ala 870 Thr aat ggg Lys Phe ggg Ala agg aag ttt Asn Asp Fer gct Ala 885 Thr aac Asn cta gat ည္တင acc Gln 900 gaa Glu g Fer Caa gcc Ala acc Thr agt Ser gat acc cta Fer gac aac Asn aca Thr Asp Thr Asp 915 Ala gat Asp gtc Val 930 ည္တ agg agt Ser aac Asn Lys 850 Gln gct Ala aac Asn tct Ser 865 cta Lea acc Thr caa



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2880	2928	2976	3024
gtt acc ttt ggc att aac acc aca agc ggt	acc cta aac gac ggt ggc ttg tct att aaa	caa atc caa gtc ggt gct gat ggc gtg aag	aat aat ggt gtt gta ggt gct ggc att gat ggc
Val Thr Phe Gly Ile Asn Thr Thr Ser Gly	Thr Leu Asn Asp Gly Gly Leu Ser Ile Lys	Gln Ile Gln Val Gly Ala Asp Gly Val Lys	Asn Asn Gly Val Val Gly Ala Gly Ile Asp Gly
955	970	990	1000
acc gac aaa aat ggt acg	ctt aaa gcc ggc aaa agc	aac ccc act ggt agc gaa	ttt gcc aag gtt aat aat
Thr Asp Lys Asn Gly Thr	Leu Lys Ala Gly Lys Ser	Asn Pro Thr Gly Ser Glu	Phe Ala Lys Val Asn Asn
945	965	980	995

3072	3120
ggg act aat ggc Gly Thr Asn Gly	yc att aac gca -y Ile Asn Ala 1040
att ggc ttt act gg Ile Gly Phe Thr Gl 1020	cta agc aaa gac ggc Leu Ser Lys Asp Gly 1035
att acc aga gat gaa Ile Thr Arg Asp Glu 1015	aaa agc aaa ccc cac Lys Ser Lys Pro His
aca act cgc a Thr Thr Arg 1	tca ctt gat a Ser Leu Asp 1



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acc aac att caa tca ggt gag att gcc caa aac Thr Asn Ile Gln Ser Gly Glu Ile Ala Gln Asn 1050 : ggt aaa aag att a 7 Gly Lys Lys Ile T 1045 ggt (

3216	3264
agc cat gat gct gtg aca ggc ggc aag att tat gat tta aaa acc gaa	ctt gaa aac aaa atc agc agt act gcc aaa aca gca caa aac tca tta
Ser His Asp Ala Val Thr Gly Gly Lys Ile Tyr Asp Leu Lys Thr Glu	Leu Glu Asn Lys Ile Ser Ser Thr Ala Lys Thr Ala Gln Asn Ser Leu
1060 1060	1075

tca gta gca gat gaa caa ggt aat aac ttt acg gtt agt Ser Val Ala Asp Glu Gln Gly Asn Asn Phe Thr Val Ser 1095

ttc

cac gaa t His Glu I 1090

3360	3408
gat gtc atc acc	: aaa ggt gtg gtg
Asp Val Ile Thr	1 Lys Gly Val Val
1120	1135
tca aag acc tct	acc aag gta aat
Ser Lys Thr Ser	Thr Lys Val Asn
1115	1130
aac cct tac tcc agt tat gac acc	t gca ggt gaa aac ggc att acc
Asn Pro Tyr Ser Ser Tyr Asp Thr	e Ala Gly Glu Asn Gly Ile Thr
1105	1125
aac Asn 1105	ttt



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3456 ogt gtg ggc att gac caa acc aaa ggc tta acc acg cct aag ctg acc Arg Val Gly Ile Asp Gln Thr Lys Gly Leu Thr Thr Pro Lys Leu Thr 1140 1150

FIG.8M

3504 ggt aat aat ggc aaa ggc att gtc att gac agc caa aat ggt Gly Asn Asn Gly Lys Gly Ile Val Ile Asp Ser Gln Asn Gly 1155 gtg Val

3552 caa aat acc atc aca gga cta agc aac act cta gct aat gtt acc aat Gln Asn Thr Ile Thr Gly Leu Ser Asn Thr Leu Ala Asn Val Thr Asn 1170 1175

3600 gat aaa ggt agc gta cgc acc aca gaa cag ggc aat ata atc aaa gac Asp Lys Gly Ser Val Arg Thr Thr Glu Gln Gly Asn Ile Ile Lys Asp 1185 3648 gaa gac aaa acc cgt gcc gcc agc att gtt gat gtg cta agc gca ggc Glu Asp Lys Thr Arg Ala Ala Ser Ile Val Asp Val Leu Ser Ala Gly 1205

3696 ttt aac ttg caa ggc aat ggt gaa gcg gtt gac ttt gtc tcc act tat Phe Asn Leu Gln Gly Asn Gly Glu Ala Val Asp Phe Val Ser Thr Tyr 1220



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3744		
g acc	l Thr	
gtg	Za.	
aag	Lys	
gct	Ala	1245
acc	Thr Ala Lys V	` '
acc	Thr	
gg	Ala	
aat	Asn	
<u>g</u>	Asp Gly Asn i	1240
gat	Asp	
gcc	Ala	
	Phe	
aac	Asn	
gtc aac	Val	1235
acc	Pr	` '
gac	Asp [

	3840
l Tyr Asp Val Asn Val 1260	a ctt ggc gta aaa acc s Leu Gly Val Lys Thr 5
Ser Lys Val Val 12	aaa gat aaa aad Lys Asp Lys Lys 1279
yar ara ayr aad arr Asp I'hr Ser Lys I'hr 1255	aca acc att gaa gtt a Thr Thr Ile Glu Val 1 1270
Tyr Asp As	gat gat ac Asp Asp Th 1265

3888		
itt gac	the Ala Leu	1295
aca ggt gct	Thr Gly Ala Asn Lys I	1290
agt act	Thr Thr Leu Thr Ser Thr Gly	1285

3936		
agt gat atc gtt gct	er Asp Ile	1310
t act ggc gat gcg ctt gtc aag gcc a	vla Leu Val Lys Ala	1305
aat caa gct act ggc gat.g	Asn Gln Ala Thr Gly Asp 7	1300

3984		
aaa ggg gca agc	Gly Ala	1325
caa act gcc	Gln Thr Ala	
t ggc gac atc	Ser Gly Asp Ile Gln Thr Ala Lys	1320
cat cta aac acc tta tc	His Leu Asn Thr Leu Se	1315



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4032	4080
tat gtg gat gct gat ggc aat aag gtc	aag tac tat caa gcc aaa aat gat ggc
Tyr Val Asp Ala Asp Gly Asn Lys Val	Lys Tyr Tyr Gln Ala Lys Asn Asp Gly
1340	1355
caa gog aac aac toa goa ggo t	atc tat gac agt acc gat aac a
Gln Ala Asn Asn Ser Ala Gly T	Ile Tyr Asp Ser Thr Asp Asn L
1330	1345

4176		
caa atg aat gtc aaa tca gtc	n Met Asn Val Lys Ser Val	1390
gct	Ala	1385
caa	Ala Gln Thr Pro Asp Gly Thr Leu	1380

gaa gtt gcc aaa gac aaa ctg gtc gcc caa Glu Val Ala Lys Asp Lys Leu Val Ala Gln 1370

: gat aaa acc aaa g . Asp Lys Thr Lys G 1365

aca gtt g Thr Val A

4224		
lat aga agg caa ggc atc	Ala Asn Ivs Ivs Gln Glv Ile Asn	1405
raa ota aat oat	Glin Glin Val Ash Ash Ala A	1400
תמט תתת בתת 11ת	Ach Tax	7395 1395

.4272		
ctt gaa aaa gcc gct tct gat	Ala Ala Ser	1420
gcc ttt gtt aaa	Ala Phe Val Lys	
gaa gac aac		1410



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ggc gtg tct ttt gta gac tca agc ggt Gly Val Ser Phe Val Asp Ser Ser Gly 1515

c aaa att gat gac aaa g r Lys Ile Asp Asp Lys G 1510

ggc acc a Gly Thr I 1505



4320	4368	4416	4464	4512
aaa acc aaa aac gcc gca gta act gtg ggt gat tta aat gcc gtt gcc	caa aca ccg ctg acc ttt gca ggg gat aca ggc aca acg gct aaa aaa	ctg ggc gag act ttg acc atc aaa ggt ggg caa aca gac acc aat aag	cta acc gat aat aac atc ggt gtg gta gca ggt act gat ggc ttc act	gtc aaa ctt gcc aaa gac cta acc aat ctt aac agc gtt aat gca ggt
Lys Thr Lys Asn Ala Ala Val Thr Val Gly Asp Leu Asn Ala Val Ala	Gln Thr Pro Leu Thr Phe Ala Gly Asp Thr Gly Thr Thr Ala Lys Lys	Leu Gly Glu Thr Leu Thr Ile Lys Gly Gly Gln Thr Asp Thr Asn Lys	Leu Thr Asp Asn Asn Ile Gly Val Val Ala Gly Thr Asp Gly Phe Thr	Val Lys Leu Ala Lys Asp Leu Thr Asn Leu Asn Ser Val Asn Ala Gly
1425 1440	1455	1460	1475 1480	1490



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4608 caa gcc aaa gca aac acc cct gtg cta agt gcc aat ggg ctg gac ctg Gln Ala Lys Ala Asn Thr Pro Val Leu Ser Ala Asn Gly Leu Asp Leu 1535 1530 1525

4656 ggc aca aaa gat acc gac Gly Gly Lys Val Ile Ser Asn Val Gly Lys Gly Thr Lys Asp Thr Asp 1550 ggt ggc aag gtc atc agt aat gtg ggc aaa 1545 1540

4704 Ala Ala Asn Val Gln Gln Leu Asn Glu Val Arg Asn Leu Leu Gly Leu caa cag tta aac gaa gta cgc aac ttg ttg ggt ctt 1565 1560 gct gcc aat gta 1555 4752 Gly Asn Ala Gly Asn Asp Asn Ala Asp Gly Asn Gln Val Asn Ile Ala ggt aat gct ggt aat gat aac gct gac ggc aat cag gta aac att gcc 1580 1575 1570 4800 1600 gac atc aaa aaa gac cca aat tca ggt tca tca tct aac cgc act gtc Asp Ile Lys Lys Asp Pro Asn Ser Gly Ser Ser Ser Asn Arg Thr Val 1595 1590

4848 atc aaa gca ggc acg gta ctt ggc ggt aaa ggt aat aac gat acc gaa Ile Lys Ala Gly Thr Val Leu Gly Gly Lys Gly Asn Asn Asp Thr Glu 1610 Ile Lys

F1G.80



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OF E JC130 SER TRADERS

4896	4944	4992	5040	2088	5136
aaa ctt gcc act ggt ggt ata caa gtg ggc gtg gat aaa gac ggc aac	gct aac ggc gat tta agc aat gtt tgg gtc aaa acc caa aaa gat ggc	agc aaa aaa gcc ctg ctc gcc act tat aac gcc gca ggt cag acc aac	tat ttg acc aac aac ccc gca gaa gcc att gac aga ata aat gaa caa	ggt atc cgc ttc ttc cat gtc aac gat ggc aat caa gag cct gtg gta	caa ggg cgt aac ggc att gac tca agt gcc tca ggc aag cac tca gtg
Lys Leu Ala Thr Gly Gly Ile Gln Val Gly Val Asp Lys Asp Gly Asn	Ala Asn Gly Asp Leu Ser Asn Val Trp Val Lys Thr Gln Lys Asp Gly	Ser Lys Lys Ala Leu Leu Ala Thr Tyr Asn Ala Ala Gly Gln Thr Asn	Tyr Leu Thr Asn Asn Pro Ala Glu Ala Ile Asp Arg Ile Asn Glu Gln	Gly Ile Arg Phe Phe His Val Asn Asp Gly Asn Gln Glu Pro Val Val	Gln Gly Arg Asn Gly Ile Asp Ser Ser Ala Ser Gly Lys His Ser Val
1620 1623	1635	1650 1650	1665 1680	1695	1710



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5184 gog ata ggt ttc cag gcc aag gca gat ggt gaa gcc gcc gtt gcc ata Ala Ile Gly Phe Gln Ala Lys Ala Asp Gly Glu Ala Ala Val Ala Ile 1715 1720

5232 ggc aga caa acc caa gca ggc aac caa tcc atc gcc atc ggt gat aac Gly Arg Gln Thr Gln Ala Gly Asn Gln Ser Ile Ala Ile Gly Asp Asn 1730

5280 gca caa gcc acg ggc gat caa tcc atc gcc atc ggt aca ggc aat gtg Ala Gln Ala Thr Gly Asp Gln Ser Ile Ala Ile Gly Thr Gly Asn Val 1745 1755 5328 gta gca ggt aag cac tct ggt gcc atc ggc gac cca agc act gtt aag Val Ala Gly Lys His Ser Gly Ala Ile Gly Asp Pro Ser Thr Val Lys 1775 5376 gct gat aac agt tac agt gtg ggt aat aac aac cag ttt acc gat gcc Ala Asp Asn Ser Tyr Ser Val Gly Asn Asn Asn Gln Phe Thr Asp Ala 1790 1785 1780

5424 act caa acc gat gtc ttt ggt gtg ggc aat aac atc acc gtg acc gaa Thr Gln Thr Asp Val Phe Gly Val Gly Asn Asn Ile Thr Val Thr Glu 1795

FIG.8S



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5472 agt aac tcg gtt gcc tta ggt tca aac tct gcc atc agt gca ggc aca Ser Asn Ser Val Ala Leu Gly Ser Asn Ser Ala Ile Ser Ala Gly Thr 1810

5520		
ggt aca	GLY	1840
gac	Asp Gly Thr	1835
aga	Lys Lys	
aca	Thr Gln Ala	1830
cac gca ggc	His Ala Gly	1825

2568		
	Ala Gly Gln	1855
ggc	Val Lys (1850
ggt gca i	Thr Ala Gly Ala Thr	1845

5616	
ggt gct gaa cgc cgt atc Gly Ala Glu Arg Arg Ile	1870
gtg ggt gcc tca ggt Val Gly Ala Ser Gly	1865
gcg gtt ggt gcg gtc tcc gl Ala Val Gly Ala Val Ser Va	1860

5664	
gcc acc agc acc gat gcg gtc Ala Thr Ser Thr Asp Ala Val	C881
ggt gag gtc agt g Gly Glu Val Ser A	1880
caa aat gtg gca gca g Gln Asn Val Ala Ala G	1875

5712		
acc caa agc att gcc aac gca acc	Thr Gln Ser Ile Ala Asn Ala	0001
ag ttg tac aaa gcc	Ser Gln Leu Tyr Lys Ala	LOOF
aat ggt agc ca	Asn Gly Ser G	



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			·		
5760	5808	5856	5904	5952	9009
aat gag ctt gac cat cgt atc cac caa aac gaa aat aag gcc aat gca Asn Glu Leu Asp His Arg Ile His Gln Asn Glu Asn Lys Ala Asn Ala 1905	ggg att tca tca gcg atg gcg atg gcg tcc atg cca caa gcc tac att Gly Ile Ser Ser Ala Met Ala Met Ala Ser Met Pro Gln Ala Tyr Ile 1935	cct ggc aga tcc atg gtt acc ggg ggt att gcc acc cac aac ggt caa Pro Gly Arg Ser Met Val Thr Gly Gly Ile Ala Thr His Asn Gly Gln 1940	ggt gcg gtg gca gtg gga ctg tcg aag ctg tcg gat aat ggt caa tgg Gly Ala Val Ala Val Gly Leu Ser Lys Leu Ser Asp Asn Gly Gln Trp 1955	gta ttt aaa atc aat ggt tca gcc gat acc caa ggc cat gta ggg gcg Val Phe Lys Ile Asn Gly Ser Ala Asp Thr Gln Gly His Val Gly Ala 1970 1970	gca gtt ggt gca ggt ttt cac ttt taagccataa atcgcaagat tttacttaaa Ala Val Gly Ala Gly Phe His Phe 1985





aatcaatete accatagttg tataaaacag catcagcate agtcatatta etgatgetga 6066

tgttttttat cacttaaacc attttaccgc tcaagtgatt ctctttcacc atgaccaaat 6126

6186 cyccattgat cataggtaaa cttattgagt aaattttatc aatgtagttg ttagatatgg

ttaaaattgt gccattgacc aaaaatgac cgatttatcc cgaaaatttc tgattatgat 6246

cogttgacct gca

6259

FIG.8V



Construction of pKS294

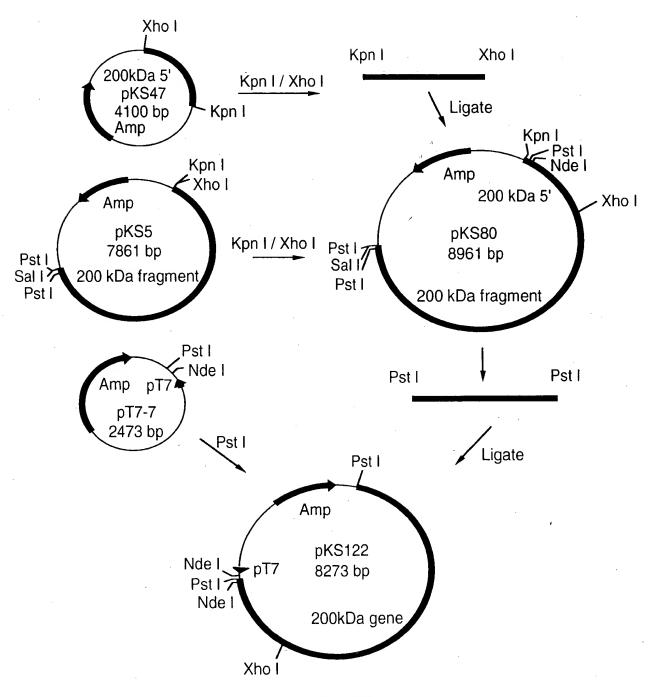


FIG.9A



Construction of pKS294

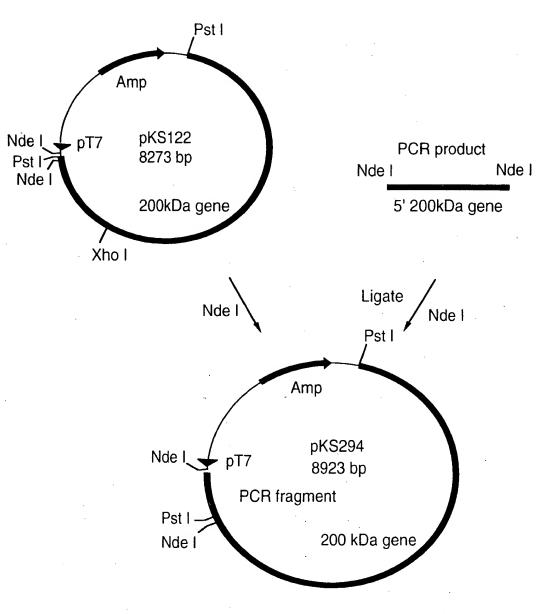


FIG.9B



Construction of pKS348

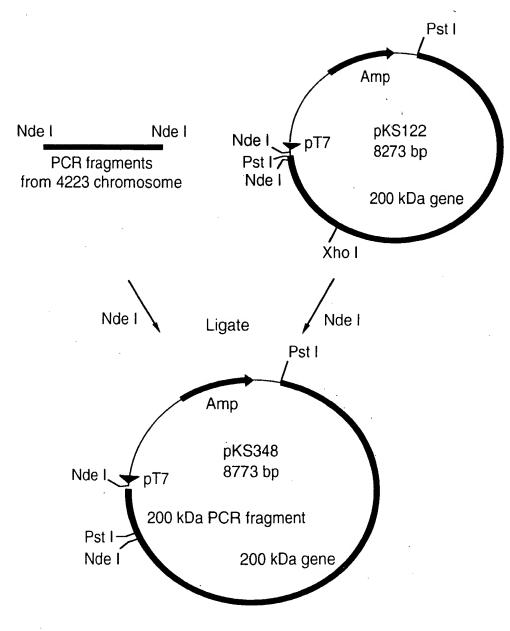


FIG.10



Purification of r200 kDa Protein from E. coli

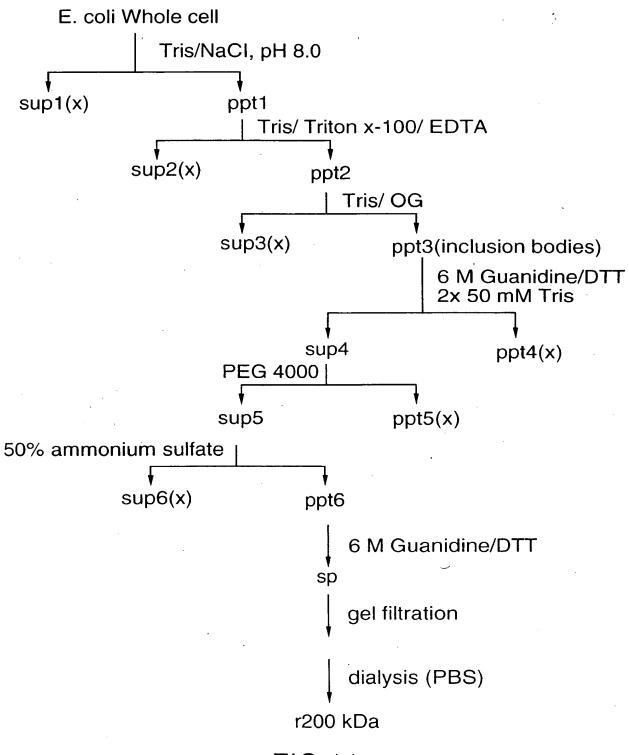
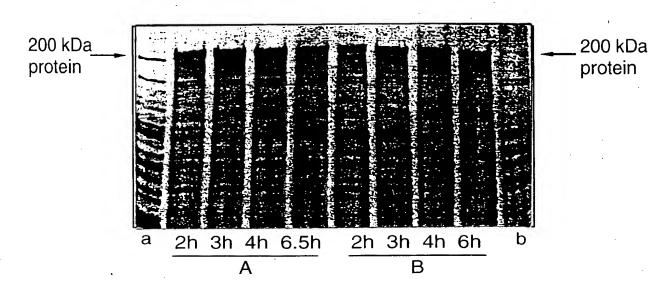


FIG.11



Expression of M 56 r200 kDa Protein Gene in E. coli



A: KS358 induced when O.D. at 600nm was 0.26

B: KS358 induced when O.D. at 600nm was 0.44

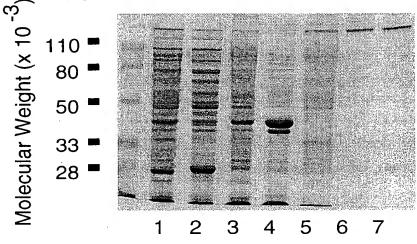
a: strain 4223 lysate

b: KS358 cultured overnight

FIG.12





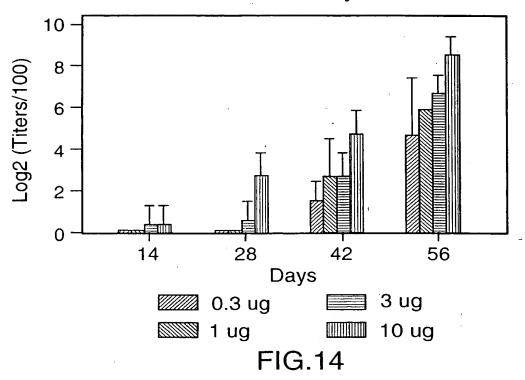


- 1. E. coli Whole cells
- 2. Soluble proteins after 50mM Tris/ NaCl, pH 8, extraction
- 3. Soluble proteins after Tris/ Triton X-100/ EDTA extraction
- 4. Soluble proteins after Tris/ OG extraction
- 5. Pellet after Tris/ OG extraction
- 6-7. Purified 200 kDa protein

FIG.13



Anti-M56 r200 kDa Antibody Titers in Mice



Anti-M56 r200 kDa Antibody Titers in Guinea Pigs

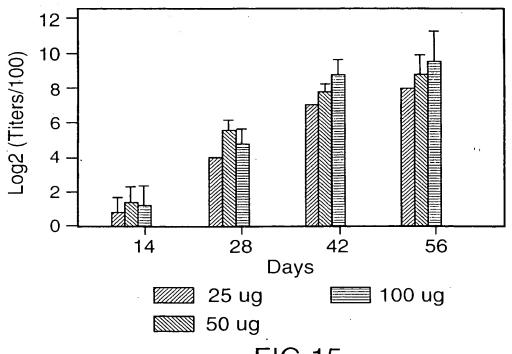


FIG.15



PCR amplification of DNA fragments carrying a portion of the 200 kDa protein gene from chromosomal DNA of RH408

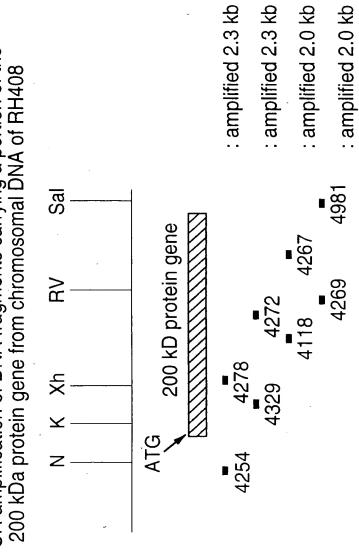


FIG.16



FIG.17A

M. catarrhalis strain 4223 200 kDa

TGCر ا ں Ç ں E ں GGTGTG Ø \mathcal{O} . G ATGGATATG ည ည

T G C C CCATACACCC Ç GGCGA Ø Ç ⊱ ø ⊱ C ِّن :

9

T G A Ø \vdash ø C E E GACAT 드 ⊱ GAT TGTAC ATC CAT

5 5 H ⊱ ď TTAAC ⊱⊣ Ø G ⊱ Ø TGTGAC Ø \mathcal{O} AA ⊱ ⊣ ⊢

TAG ⊱⊣ [---⋖ Ø \vdash Ø \mathcal{C} TTAC ø \mathcal{O} [--1 Ø \mathcal{C} \mathcal{O} T G G ⊣ C Ø \vdash AA

140

150

AAAAAT ⊱ E G TTAGTAA ΑT $^{\circ}$ G A C ø ⊱

180 CGCATT? 160

T G A A 210 Ø ⊱ K ں ⊱ TGTA T G Ø ⊱⊣ ⊱ ည ည G C C T G C Ø

190



FIG. 17B

G T ⊱ ⊏ Ø ⊣ H Ø G $T \subset T$ Ø ⊣ G \vdash ₽ ø Ç ⊱ T T ⊱ Ø ⊱ Ø G K ⊱⊣

220

Ċ ⊱ K \mathcal{O} E Ø G \vdash TA ø ⊱ \mathcal{O} G \vdash Ø \mathcal{O} \vdash G G A T 250 Ø G Ø \mathcal{O} Ø

260

d 亡 ں ⊱ ں ø ⊣ AA 290 Н ⊱ G c C \vdash 드 G A 든 G ø Ç \mathcal{O} ₽

G ⊢ \vdash Ø ⊱ \mathcal{O} ш Ø K ר ט Ø K Ø Ç E \vdash E⊣ \vdash Ø E TA K Ç ⊱ [-- ₽ Ø A G E ⊱ ⊱ ď Ø ⊱ ΤA ⊱ Ç G \vdash ø ⊣ Ø \mathcal{O} A C J E Ø Ø K E ⊱

360

Ø \mathcal{O} K \mathcal{O} 든 Ø Ø ø A A 380 ⊱⊣ \mathcal{O} ⊱⊣ 드 ⊱ E-- \vdash \mathcal{O} G G T A 370 G G H K K \vdash

E G Ø \mathcal{O} \mathcal{O} K TA(G \vdash ى Ø \vdash \mathcal{O} 든 G T ⊱ ⊱ K ⊱ Ø Ø \mathcal{O} \mathcal{O} 420

450 Ø \mathcal{O} ⊱ Ø A G ں A C ں ⊱ Ø Ø \mathcal{O} \mathcal{O} \circ ⊱ K Ç A C 430 Ċ ⊣ K Ø Ç ⊣



FIG.17C

... CAAGTGATGTTTTGTATACGCACCATTTA

460 470

CCCTAATTTTCAATCAATGCCTATGTC

500 510

... AGCATGTATCATTTTTTTAAGGTAAACCAC

MET ASN HIS ILE TYR LYS VAL ILE PHE ASN

CATGAATCACATCTATAAAGTCATCTTAA

260

LYS ALA 12 THR GLY THR PHE MET ALI

 $\dot{\rm VAL}^{19}$ ALA

...CAAAGCCACAGACATTTATGGCAGTGGC

580 590

GLU TYR ALA LYS SER HIS SER THR GLY GLY AGAGTACGCCAAATCCCACAGCACGGGGG 620 630



...GGGTAGCTGCTACAGGGCAAGTTGGCAG S S 650 铝 640 SE

GLY39 SER

999

TGTATGCACTCTGAGCTTTGCCCGTATTGC 9 ALA 出

ALA VAL 昌

...CGCGCTCTGTCCTCGTGATCGGTGCAAC

FIG.17D



3' Half Constructs Of 200 kD Protein Gene

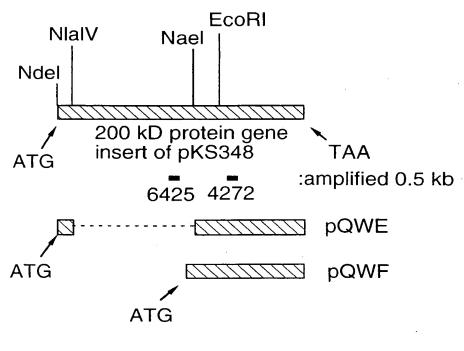


FIG.18



Construction of pQWE

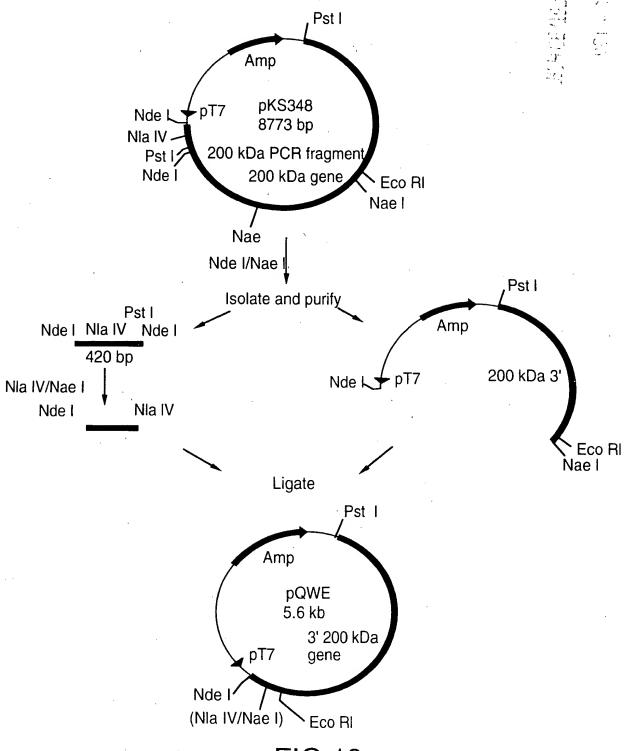


FIG.19



